GenCore version 5.1.8

OM protein - protein search, using sw model

Run on: May 13, 2006, 08:14:12 ; Search time 46 Seconds
(without alignments)
25.162 Million cell updates/sec

Title: US-10-769-514-17
Perfect score: 74
Sequence: 1 MGYGMALSKINLHN 14

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 100 summaries

Database : Issued_Patents_AA:*
1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/6_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	74	100.0	682	2	US-08-613-009A-10	Sequence 10, Appl
2	74	100.0	682	2	US-08-778-570B-12	Sequence 12, Appl
3	74	100.0	682	2	US-09-059-584-12	Sequence 12, Appl
4	74	100.0	702	1	US-08-867-941-25	Sequence 25, Appl
5	74	100.0	702	2	US-08-613-009A-9	Sequence 9, Appli
6	74	100.0	702	2	US-09-074-658-25	Sequence 25, Appl
7	74	100.0	702	2	US-08-778-570B-11	Sequence 11, Appl
8	74	100.0	702	2	US-09-059-584-11	Sequence 11, Appl
9	65	87.8	706	2	US-09-059-584-46	Sequence 46, Appl
10	63	85.1	713	2	US-09-059-584-49	Sequence 49, Appl
11	42	56.8	696	2	US-09-907-794A-91	Sequence 91, Appl
12	42	56.8	696	2	US-09-905-125A-91	Sequence 91, Appl
13	42	56.8	696	2	US-09-902-775A-91	Sequence 91, Appl
14	42	56.8	696	2	US-09-906-700-91	Sequence 91, Appl
15	42	56.8	696	2	US-09-903-603A-91	Sequence 91, Appl
16	42	56.8	696	2	US-09-904-920A-91	Sequence 91, Appl
17	42	56.8	696	2	US-09-909-064-91	Sequence 91, Appl
18	42	56.8	696	2	US-09-905-381A-91	Sequence 91, Appl
19	42	56.8	696	2	US-09-906-618-91	Sequence 91, Appl
20	42	56.8	696	2	US-09-906-646-91	Sequence 91, Appl
21	42	56.8	696	2	US-09-904-462-91	Sequence 91, Appl
22	42	56.8	696	2	US-09-902-736A-91	Sequence 91, Appl
23	42	56.8	696	2	US-09-906-722A-91	Sequence 91, Appl
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25	40	54.1	81	2	US-09-270-767-54638	Sequence 54638, A
26	40	54.1	622	2	US-09-902-540-11017	Sequence 11017, A
27	39	52.7	315	2	US-09-134-000C-3657	Sequence 3657, Ap
28	38	51.4	130	2	US-09-328-352-7870	Sequence 7870, Ap
29	38	51.4	282	2	US-09-561-077C-32	Sequence 32, Appl
30	38	51.4	282	2	US-09-221-014-32	Sequence 32, Appl
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32	37.5	50.7	596	2	US-09-631-022-6	Sequence 6, Appli
33	37	50.0	42	2	US-09-830-807-7	Sequence 7, Appli
34	37	50.0	167	2	US-09-270-767-37334	Sequence 37334, A
35	37	50.0	167	2	US-09-270-767-52551	Sequence 52551, A
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37	37	50.0	228	2	US-09-949-016-9719	Sequence 9719, Ap
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42	36	48.6	192	2	US-09-252-991A-21776	Sequence 21776, A
43	36	48.6	305	2	US-09-315-794-22	Sequence 22, Appl
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45	36	48.6	310	2	US-09-107-532A-6246	Sequence 6246, Ap
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47	36	48.6	486	2	US-09-902-540-10512	Sequence 10512, A
48	36	48.6	854	2	US-10-335-711-9	Sequence 9, Appli
49	36	48.6	1115	2	US-10-335-711-2	Sequence 2, Appli
50	36	48.6	1564	2	US-09-976-594-309	Sequence 309, App
51	36	48.6	2763	2	US-08-496-944-2	Sequence 2, Appli
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55	35	47.3	150	2	US-09-270-767-48753	Sequence 48753, A
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59	35	47.3	151	2	US-09-270-767-48435	Sequence 48435, A
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61	35	47.3	152	2	US-09-270-767-47371	Sequence 47371, A
62	35	47.3	179	2	US-09-270-767-33076	Sequence 33076, A
63	35	47.3	179	2	US-09-270-767-48293	Sequence 48293, A
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66	35	47.3	370	1	US-08-729-214-27	Sequence 27, Appl
67	35	47.3	370	2	US-09-028-934-27	Sequence 27, Appl
68	35	47.3	373	2	US-09-028-934-32	Sequence 32, Appl
69	35	47.3	407	2	US-08-732-749-2	Sequence 2, Appli
70	35	47.3	407	2	US-09-258-613-1	Sequence 1, Appli
71	35	47.3	407	2	US-10-156-300-1	Sequence 1, Appli
72	35	47.3	407	2	US-09-950-739-2	Sequence 2, Appli
73	35	47.3	408	2	US-09-270-767-32356	Sequence 32356, A
74	35	47.3	408	2	US-09-270-767-47573	Sequence 47573, A
75	35	47.3	631	2	US-09-949-016-10729	Sequence 10729, A
76	35	47.3	688	2	US-09-762-767B-2	Sequence 2, Appli
77	35	47.3	717	2	US-09-949-016-5999	Sequence 5999, Ap
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79	35	47.3	1259	2	US-09-489-039A-8840	Sequence 8840, Ap
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81	35	47.3	1498	1	US-08-404-531B-29	Sequence 29, Appl
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84	35	47.3	1498	2	US-08-488-546A-28	Sequence 28, Appl
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86	35	47.3	1528	1	US-08-463-092B-6	Sequence 6, Appli
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88	35	47.3	1528	1	US-08-460-907B-6	Sequence 6, Appli
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92	35	47.3	1580	2	US-09-208-716-1	Sequence 1, Appli
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96	35	47.3	1581	2	US-08-726-320-3	Sequence 3, Appli
97	35	47.3	1581	2	US-08-726-320-4	Sequence 4, Appli
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99	35	47.3	1581	2	US-09-208-716-4	Sequence 4, Appli
100	35	47.3	1582	1	US-08-404-531B-9	Sequence 9, Appli

ALIGNMENTS

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RESULT 1
US-08-613-009A-10
; Sequence 10, Application US/08613009A
; Patent No. 6090576
;  GENERAL INFORMATION:
;    APPLICANT:  Myers, Lisa E
;    APPLICANT:  Schryvers, Anthony B
;    APPLICANT:  Harkness, Robin E
;    APPLICANT:  Loosmore, Sheena M.
;    APPLICANT:  Du, Run-Pan
;    APPLICANT:  Yang, Yan-Ping
;    APPLICANT:  Klein, Michel H
;    TITLE OF INVENTION:  Transferrin Receptor Genes of Moraxella
;    NUMBER OF SEQUENCES:  31
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Sim & McBurney
;      STREET:  6th Floor, 330 University Avenue
;      CITY:  Toronto
;      STATE:  Ontario
;      COUNTRY:  Canada
;      ZIP:  M5G 1R7
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/613,009A
;      FILING DATE:  08-MAR-1996
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Stewart, Michael I
;      REGISTRATION NUMBER:  24973
;      REFERENCE/DOCKET NUMBER:  1038-542
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (416) 595-1155
;      TELEFAX:  (416) 595-1163
;  INFORMATION FOR SEQ ID NO:  10:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  682 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-613-009A-10

  Query Match             100.0%;  Score 74;  DB 2;  Length 682;
  Best Local Similarity   100.0%;  Pred. No. 1.9e-05;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
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Db         80 MGYGMALSKINLHN 93


RESULT 2
US-08-778-570B-12
; Sequence 12, Application US/08778570B
; Patent No. 6437096
;  GENERAL INFORMATION:
;    APPLICANT:  Myers, Lisa E
;    APPLICANT:  Schryvers, Anthony B
;    APPLICANT:  Harkness, Robin E
;    APPLICANT:  Loosmore, Sheena M.
;    APPLICANT:  Du, Run-Pan
;    APPLICANT:  Yang, Yan-Ping
;    APPLICANT:  Klein, Michel H
;    TITLE OF INVENTION:  Transferrin Receptor Genes of Moraxella
;    NUMBER OF SEQUENCES:  43
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Sim & McBurney
;      STREET:  6th Floor, 330 University Avenue
;      CITY:  Toronto
;      STATE:  Ontario
;      COUNTRY:  Canada
;      ZIP:  M5G 1R7
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/778,570B
;      FILING DATE:  03-JAN-1997
;      CLASSIFICATION:  536
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Stewart, Michael I
;      REGISTRATION NUMBER:  24973
;      REFERENCE/DOCKET NUMBER:  1038-664
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (416) 595-1155
;      TELEFAX:  (416) 595-1163
;  INFORMATION FOR SEQ ID NO:  12:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  682 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-778-570B-12

  Query Match             100.0%;  Score 74;  DB 2;  Length 682;
  Best Local Similarity   100.0%;  Pred. No. 1.9e-05;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||||
Db         80 MGYGMALSKINLHN 93


RESULT 3
US-09-059-584-12
; Sequence 12, Application US/09059584
; Patent No. 6440701
;  GENERAL INFORMATION:
;    APPLICANT:  Myers, Lisa E
;    APPLICANT:  Schryvers, Anthony B
;    APPLICANT:  Harkness, Robin E
;    APPLICANT:  Loosmore, Sheena M.
;    APPLICANT:  Du, Run-Pan
;    APPLICANT:  Yang, Yan-Ping
;    APPLICANT:  Klein, Michel H
;    TITLE OF INVENTION:  Transferrin Receptor Genes of Moraxella
;    NUMBER OF SEQUENCES:  60
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Sim & McBurney
;      STREET:  6th Floor, 330 University Avenue
;      CITY:  Toronto
;      STATE:  Ontario
;      COUNTRY:  Canada
;      ZIP:  M5G 1R7
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/059,584
;      FILING DATE:  14-APR-1998
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/778,570
;      FILING DATE:  03-JAN-1997
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;      CLASSIFICATION:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Stewart, Michael I
;      REGISTRATION NUMBER:  24973
;      REFERENCE/DOCKET NUMBER:  1038-794
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (416) 595-1155
;      TELEFAX:  (416) 595-1163
;  INFORMATION FOR SEQ ID NO:  12:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  682 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-09-059-584-12

  Query Match             100.0%;  Score 74;  DB 2;  Length 682;
  Best Local Similarity   100.0%;  Pred. No. 1.9e-05;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
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Db         80 MGYGMALSKINLHN 93


RESULT 4
US-08-867-941-25
; Sequence 25, Application US/08867941
; Patent No. 5977337
;  GENERAL INFORMATION:
;    APPLICANT:  Loosmore, Sheena M
;    APPLICANT:  Du, Run-Pan
;    APPLICANT:  Wang, Quijun
;    APPLICANT:  Yang, Yan-Ping
;    APPLICANT:  Klein, Michel H
;    TITLE OF INVENTION:  LACTOFERRIN RECEPTOR GENES OF MORAXELLA
;    NUMBER OF SEQUENCES:  67
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Sim & McBurney
;      STREET:  6th Floor, 330 University Avenue
;      CITY:  Toronto
;      STATE:  Ontario
;      COUNTRY:  Canada
;      ZIP:  M5G 1R7
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/867,941
;      FILING DATE:  03-JUN-1997
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Stewart, Michael I
;      REGISTRATION NUMBER:  24,973
;      REFERENCE/DOCKET NUMBER:  1038-681 MIS:jb
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (416) 595-1155
;      TELEFAX:  (416) 595-1163
;  INFORMATION FOR SEQ ID NO:  25:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  702 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-867-941-25

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  Best Local Similarity   100.0%;  Pred. No. 2e-05;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
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Db        100 MGYGMALSKINLHN 113


RESULT 5
US-08-613-009A-9
; Sequence 9, Application US/08613009A
; Patent No. 6090576
;  GENERAL INFORMATION:
;    APPLICANT:  Myers, Lisa E
;    APPLICANT:  Schryvers, Anthony B
;    APPLICANT:  Harkness, Robin E
;    APPLICANT:  Loosmore, Sheena M.
;    APPLICANT:  Du, Run-Pan
;    APPLICANT:  Yang, Yan-Ping
;    APPLICANT:  Klein, Michel H
;    TITLE OF INVENTION:  Transferrin Receptor Genes of Moraxella
;    NUMBER OF SEQUENCES:  31
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Sim & McBurney
;      STREET:  6th Floor, 330 University Avenue
;      CITY:  Toronto
;      STATE:  Ontario
;      COUNTRY:  Canada
;      ZIP:  M5G 1R7
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/613,009A
;      FILING DATE:  08-MAR-1996
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Stewart, Michael I
;      REGISTRATION NUMBER:  24973
;      REFERENCE/DOCKET NUMBER:  1038-542
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (416) 595-1155
;      TELEFAX:  (416) 595-1163
;  INFORMATION FOR SEQ ID NO:  9:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  702 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-613-009A-9

  Query Match             100.0%;  Score 74;  DB 2;  Length 702;
  Best Local Similarity   100.0%;  Pred. No. 2e-05;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||||
Db        100 MGYGMALSKINLHN 113


RESULT 6
US-09-074-658-25
; Sequence 25, Application US/09074658
; Patent No. 6184371
;  GENERAL INFORMATION:
;    APPLICANT:  Loosmore, Sheena M
;    APPLICANT:  Run-Pan Du
;    APPLICANT:  Quijun Wang
;    APPLICANT:  Yang, Yan-Ping
;    APPLICANT:  Klein, Michel H
;    TITLE OF INVENTION:  LACTOFERRIN RECEPTOR GENES OF MORAXELLA
;    NUMBER OF SEQUENCES:  78
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Sim & McBurney
;      STREET:  6th Floor, 330 University Avenue
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;      CITY:  Toronto
;      STATE:  Ontario
;      COUNTRY:  Canada
;      ZIP:  M5G 1R7
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/074,658
;      FILING DATE:  08-MAY-1998
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Stewart, Michael I
;      REGISTRATION NUMBER:  24,973
;      REFERENCE/DOCKET NUMBER:  1038-795
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (416) 595-1155
;      TELEFAX:  (416) 595-1163
;  INFORMATION FOR SEQ ID NO:  25:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  702 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-09-074-658-25

  Query Match             100.0%;  Score 74;  DB 2;  Length 702;
  Best Local Similarity   100.0%;  Pred. No. 2e-05;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||||
Db        100 MGYGMALSKINLHN 113


RESULT 7
US-08-778-570B-11
; Sequence 11, Application US/08778570B
; Patent No. 6437096
;  GENERAL INFORMATION:
;    APPLICANT:  Myers, Lisa E
;    APPLICANT:  Schryvers, Anthony B
;    APPLICANT:  Harkness, Robin E
;    APPLICANT:  Loosmore, Sheena M.
;    APPLICANT:  Du, Run-Pan
;    APPLICANT:  Yang, Yan-Ping
;    APPLICANT:  Klein, Michel H
;    TITLE OF INVENTION:  Transferrin Receptor Genes of Moraxella
;    NUMBER OF SEQUENCES:  43
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Sim & McBurney
;      STREET:  6th Floor, 330 University Avenue
;      CITY:  Toronto
;      STATE:  Ontario
;      COUNTRY:  Canada
;      ZIP:  M5G 1R7
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/778,570B
;      FILING DATE:  03-JAN-1997
;      CLASSIFICATION:  536
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Stewart, Michael I
;      REGISTRATION NUMBER:  24973
;      REFERENCE/DOCKET NUMBER:  1038-664
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (416) 595-1155
;      TELEFAX:  (416) 595-1163
;  INFORMATION FOR SEQ ID NO:  11:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  702 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-778-570B-11

  Query Match             100.0%;  Score 74;  DB 2;  Length 702;
  Best Local Similarity   100.0%;  Pred. No. 2e-05;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||||
Db        100 MGYGMALSKINLHN 113


RESULT 8
US-09-059-584-11
; Sequence 11, Application US/09059584
; Patent No. 6440701
;  GENERAL INFORMATION:
;    APPLICANT:  Myers, Lisa E
;    APPLICANT:  Schryvers, Anthony B
;    APPLICANT:  Harkness, Robin E
;    APPLICANT:  Loosmore, Sheena M.
;    APPLICANT:  Du, Run-Pan
;    APPLICANT:  Yang, Yan-Ping
;    APPLICANT:  Klein, Michel H
;    TITLE OF INVENTION:  Transferrin Receptor Genes of Moraxella
;    NUMBER OF SEQUENCES:  60
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Sim & McBurney
;      STREET:  6th Floor, 330 University Avenue
;      CITY:  Toronto
;      STATE:  Ontario
;      COUNTRY:  Canada
;      ZIP:  M5G 1R7
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/059,584
;      FILING DATE:  14-APR-1998
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/778,570
;      FILING DATE:  03-JAN-1997
;      CLASSIFICATION:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Stewart, Michael I
;      REGISTRATION NUMBER:  24973
;      REFERENCE/DOCKET NUMBER:  1038-794
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (416) 595-1155
;      TELEFAX:  (416) 595-1163
;  INFORMATION FOR SEQ ID NO:  11:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  702 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-09-059-584-11

  Query Match             100.0%;  Score 74;  DB 2;  Length 702;
  Best Local Similarity   100.0%;  Pred. No. 2e-05;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||||
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Db        100 MGYGMALSKINLHN 113


RESULT 9
US-09-059-584-46
; Sequence 46, Application US/09059584
; Patent No. 6440701
;  GENERAL INFORMATION:
;    APPLICANT:  Myers, Lisa E
;    APPLICANT:  Schryvers, Anthony B
;    APPLICANT:  Harkness, Robin E
;    APPLICANT:  Loosmore, Sheena M.
;    APPLICANT:  Du, Run-Pan
;    APPLICANT:  Yang, Yan-Ping
;    APPLICANT:  Klein, Michel H
;    TITLE OF INVENTION:  Transferrin Receptor Genes of Moraxella
;    NUMBER OF SEQUENCES:  60
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Sim & McBurney
;      STREET:  6th Floor, 330 University Avenue
;      CITY:  Toronto
;      STATE:  Ontario
;      COUNTRY:  Canada
;      ZIP:  M5G 1R7
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/059,584
;      FILING DATE:  14-APR-1998
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/778,570
;      FILING DATE:  03-JAN-1997
;      CLASSIFICATION:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Stewart, Michael I
;      REGISTRATION NUMBER:  24973
;      REFERENCE/DOCKET NUMBER:  1038-794
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (416) 595-1155
;      TELEFAX:  (416) 595-1163
;  INFORMATION FOR SEQ ID NO:  46:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  706 amino acids
;      TYPE:  amino acid
;      TOPOLOGY:  linear
US-09-059-584-46

  Query Match             87.8%;  Score 65;  DB 2;  Length 706;
  Best Local Similarity   92.3%;  Pred. No. 0.00097;
  Matches   12;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLH 13
              |||||||||:|||
Db        100 MGYGMALSKMNLH 112


RESULT 10
US-09-059-584-49
; Sequence 49, Application US/09059584
; Patent No. 6440701
;  GENERAL INFORMATION:
;    APPLICANT:  Myers, Lisa E
;    APPLICANT:  Schryvers, Anthony B
;    APPLICANT:  Harkness, Robin E
;    APPLICANT:  Loosmore, Sheena M.
;    APPLICANT:  Du, Run-Pan
;    APPLICANT:  Yang, Yan-Ping
;    APPLICANT:  Klein, Michel H
;    TITLE OF INVENTION:  Transferrin Receptor Genes of Moraxella
;    NUMBER OF SEQUENCES:  60
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Sim & McBurney
;      STREET:  6th Floor, 330 University Avenue
;      CITY:  Toronto
;      STATE:  Ontario
;      COUNTRY:  Canada
;      ZIP:  M5G 1R7
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/059,584
;      FILING DATE:  14-APR-1998
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/778,570
;      FILING DATE:  03-JAN-1997
;      CLASSIFICATION:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Stewart, Michael I
;      REGISTRATION NUMBER:  24973
;      REFERENCE/DOCKET NUMBER:  1038-794
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (416) 595-1155
;      TELEFAX:  (416) 595-1163
;  INFORMATION FOR SEQ ID NO:  49:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  713 amino acids
;      TYPE:  amino acid
;      TOPOLOGY:  linear
US-09-059-584-49

  Query Match             85.1%;  Score 63;  DB 2;  Length 713;
  Best Local Similarity   85.7%;  Pred. No. 0.0023;
  Matches   12;  Conservative    2;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||::
Db        100 MGYGMALSKINLYD 113


RESULT 11
US-09-907-794A-91
; Sequence 91, Application US/09907794A
; Patent No. 6635468
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumás, Daniel
;  APPLICANT:  Williams, P. Mickey
```

```
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/907,794A
;  CURRENT FILING DATE:  2001-07-17
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;    LENGTH: 696
;    TYPE: PRT
;    ORGANISM: Homo sapiens
US-09-907-794A-91

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 20;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 12
US-09-905-125A-91
; Sequence 91, Application US/09905125A
; Patent No. 6664376
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/905,125A
;  CURRENT FILING DATE:  2001-07-12
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;    LENGTH: 696
;    TYPE: PRT
;    ORGANISM: Homo sapiens
US-09-905-125A-91

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 20;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 13
US-09-902-775A-91
; Sequence 91, Application US/09902775A
; Patent No. 6686451
; GENERAL INFORMATION:
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; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/902,775A
; CURRENT FILING DATE: 2001-07-10
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-902-775A-91

Query Match 56.8%; Score 42; DB 2; Length 696;
Best Local Similarity 63.6%; Pred. No. 20;

Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||
Db 493 GVSLSKLSLHN 503

RESULT 14
US-09-906-700-91
; Sequence 91, Application US/09906700
; Patent No. 6723535
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/906,700
; CURRENT FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999

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;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-906-700-91

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 20;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 15
US-09-903-603A-91
; Sequence 91, Application US/09903603A
; Patent No. 6767995
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: GNE.1618P2C12
;  CURRENT APPLICATION NUMBER: US/09/903,603A
;  CURRENT FILING DATE:  2001-07-11
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-903-603A-91

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 20;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 16
US-09-904-920A-91
; Sequence 91, Application US/09904920A
; Patent No. 6806352
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,920A
;  CURRENT FILING DATE:  2001-07-13
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
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;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-904-920A-91

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 20;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 17
US-09-909-064-91
; Sequence 91, Application US/09909064
; Patent No. 6818449
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/909,064
;  CURRENT FILING DATE:  2001-07-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-909-064-91

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 20;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 18
US-09-905-381A-91
; Sequence 91, Application US/09905381A
; Patent No. 6818746
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
```

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;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/905,381A
;  CURRENT FILING DATE:  2001-07-13
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-905-381A-91

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 20;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |:|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 19
US-09-906-618-91
; Sequence 91, Application US/09906618
; Patent No. 6828146
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,618
;  CURRENT FILING DATE:  2001-07-16
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-906-618-91

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 20;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;
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Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 20
US-09-906-646-91
; Sequence 91, Application US/09906646
; Patent No. 6852848
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,646
;  CURRENT FILING DATE:  2002-01-22
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-906-646-91

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 20;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 21
US-09-904-462-91
; Sequence 91, Application US/09904462
; Patent No. 6878807
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,462
;  CURRENT FILING DATE:  2001-07-13
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
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; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-904-462-91

Query Match 56.8%; Score 42; DB 2; Length 696;
Best Local Similarity 63.6%; Pred. No. 20;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

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Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
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RESULT 22
US-09-902-736A-91
; Sequence 91, Application US/09902736A
; Patent No. 6894148
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/902,736A
; CURRENT FILING DATE: 2001-07-10
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-902-736A-91

Query Match 56.8%; Score 42; DB 2; Length 696;
Best Local Similarity 63.6%; Pred. No. 20;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

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Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 23
US-09-906-722A-91
; Sequence 91, Application US/09906722A
; Patent No. 6946262
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey

```
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: GNE.1618P2C61
;  CURRENT APPLICATION NUMBER: US/09/906,722A
;  CURRENT FILING DATE:  2001-07-16
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-906-722A-91

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 20;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 24
US-09-270-767-39421
; Sequence 39421, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
;  APPLICANT: Homburger et al.
;  TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
;  FILE REFERENCE: File Reference: 7326-094
;  CURRENT APPLICATION NUMBER: US/09/270,767
;  CURRENT FILING DATE: 1999-03-17
;  NUMBER OF SEQ ID NOS: 62517
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 39421
;   LENGTH: 81
;   TYPE: PRT
;   ORGANISM: Drosophila melanogaster
;   FEATURE:
;   OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-39421

  Query Match             54.1%;  Score 40;  DB 2;  Length 81;
  Best Local Similarity   53.8%;  Pred. No. 4.1;
  Matches    7;  Conservative    3;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              ||  :: :|||||
Db         45 GYMHSIPRINLHN 57


RESULT 25
US-09-270-767-54638
; Sequence 54638, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
;  APPLICANT: Homburger et al.
;  TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
;  FILE REFERENCE: File Reference: 7326-094
;  CURRENT APPLICATION NUMBER: US/09/270,767
;  CURRENT FILING DATE: 1999-03-17
;  NUMBER OF SEQ ID NOS: 62517
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 54638
;   LENGTH: 81
;   TYPE: PRT
;   ORGANISM: Drosophila melanogaster
;   FEATURE:
;   OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-54638

  Query Match             54.1%;  Score 40;  DB 2;  Length 81;
  Best Local Similarity   53.8%;  Pred. No. 4.1;
  Matches    7;  Conservative    3;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              ||  :: :|||||
Db         45 GYMHSIPRINLHN 57


RESULT 26
US-09-902-540-11017
; Sequence 11017, Application US/09902540
; Patent No. 6833447
; GENERAL INFORMATION:
;  APPLICANT: Goldman, Barry S.
;  APPLICANT:  Hinkle, Gregory J.
;  APPLICANT:  Slater, Steven C.
;  APPLICANT:  Wiegand, Roger C.
;  TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof
;  FILE REFERENCE: 38-10(15849)B
;  CURRENT APPLICATION NUMBER: US/09/902,540
;  CURRENT FILING DATE:  2001-07-10
;  PRIOR APPLICATION NUMBER: 60/217,883
;  PRIOR FILING DATE: 2000-07-10
;  NUMBER OF SEQ ID NOS: 16825
; SEQ ID NO 11017
;   LENGTH: 622
;   TYPE: PRT
;   ORGANISM: Myxococcus xanthus
US-09-902-540-11017

  Query Match             54.1%;  Score 40;  DB 2;  Length 622;
  Best Local Similarity   63.6%;  Pred. No. 42;
  Matches    7;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINL 12
              |||| ||:::|
Db        405 GYGMLLSRLSL 415


RESULT 27
```

```
US-09-134-000C-3657
; Sequence 3657, Application US/09134000C
; Patent No. 6617156
; GENERAL INFORMATION:
;  APPLICANT: Lynn Doucette-Stamm et al
;  TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
;  TITLE OF INVENTION:  ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS
;  FILE REFERENCE: 032796-032
;  CURRENT APPLICATION NUMBER: US/09/134,000C
;  CURRENT FILING DATE:  1998-08-13
;  PRIOR APPLICATION NUMBER: US 60/055,778
;  PRIOR FILING DATE: 1997-08-15
;  NUMBER OF SEQ ID NOS: 6812
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3657
;   LENGTH: 315
;   TYPE: PRT
;   ORGANISM: Enterococcus faecalis
US-09-134-000C-3657

  Query Match             52.7%;  Score 39;  DB 2;  Length 315;
  Best Local Similarity   54.5%;  Pred. No. 30;
  Matches    6;  Conservative    4;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINL 12
              |:|:|::| ||
Db         25 GFGLAITKFNL 35


RESULT 28
US-09-328-352-7870
; Sequence 7870, Application US/09328352
; Patent No. 6562958
; GENERAL INFORMATION:
;  APPLICANT: Gary L. Breton et al.
;  TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
;  TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
;  FILE REFERENCE: GTC99-03PA
;  CURRENT APPLICATION NUMBER: US/09/328,352
;  CURRENT FILING DATE: 1999-06-04
;  NUMBER OF SEQ ID NOS: 8252
; SEQ ID NO 7870
;   LENGTH: 130
;   TYPE: PRT
;   ORGANISM: Acinetobacter baumannii
US-09-328-352-7870

  Query Match             51.4%;  Score 38;  DB 2;  Length 130;
  Best Local Similarity   50.0%;  Pred. No. 17;
  Matches    6;  Conservative    5;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          3 YGMALSKINLHN 14
              :|||:| |::|:
Db         99 WGMAMSAISIHS 110


RESULT 29
US-09-561-077C-32
; Sequence 32, Application US/09561077C
; Patent No. 6706501
; GENERAL INFORMATION:
;  APPLICANT: Rosson, Reinhardt D.
;  APPLICANT:  Deng, Ming-de
;  APPLICANT:  Grund, Alan D.
;  TITLE OF INVENTION: LINOLEATE ISOMERASE
;  FILE REFERENCE: 3161-20-C1
;  CURRENT APPLICATION NUMBER: US/09/561,077C
;  CURRENT FILING DATE:  2000-04-28
;  PRIOR APPLICATION NUMBER: 60/141,798
;  PRIOR FILING DATE: 1999-06-30
;  NUMBER OF SEQ ID NOS: 80
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 32
;   LENGTH: 282
;   TYPE: PRT
;   ORGANISM: Lactobacillus reuteri
US-09-561-077C-32

  Query Match             51.4%;  Score 38;  DB 2;  Length 282;
  Best Local Similarity   46.2%;  Pred. No. 41;
  Matches    6;  Conservative    4;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLH 13
              : ||: |:|: ||
Db        132 LNYGIWLNKVRLH 144


RESULT 30
US-09-221-014-32
; Sequence 32, Application US/09221014C
; Patent No. 6743609
; GENERAL INFORMATION:
;  APPLICANT: Rosson, Reinhardt D.
;  APPLICANT: Grund, Alan D.
;  APPLICANT: Deng, Ming-de
;  APPLICANT: Sanchez-Riera, Fernando
;  TITLE OF INVENTION: LINOLEATE ISOMERASE
;  FILE REFERENCE: 3161-20
;  CURRENT APPLICATION NUMBER: US/09/221,014C
;  CURRENT FILING DATE: 1998-12-23
;  EARLIER APPLICATION NUMBER: 60/068,617
;  EARLIER FILING DATE: 1997-12-23
;  EARLIER APPLICATION NUMBER: 60/089,560
;  EARLIER FILING DATE: 1998-06-17
;  NUMBER OF SEQ ID NOS: 41
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 32
;   LENGTH: 282
;   TYPE: PRT
;   ORGANISM: Lactobacillus reuteri
US-09-221-014-32

  Query Match             51.4%;  Score 38;  DB 2;  Length 282;
  Best Local Similarity   46.2%;  Pred. No. 41;
  Matches    6;  Conservative    4;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLH 13
              : ||: |:|: ||
Db        132 LNYGIWLNKVRLH 144


RESULT 31
US-09-171-337A-6
; Sequence 6, Application US/09171337A
; Patent No. 6300095
;   GENERAL INFORMATION:
;        APPLICANT: BARREDO FUENTE, Jose Luis
;                   RODRIGUEZ SAIZ, Marta
;                   COLLADOS DE LA VIEJA, Alfonso J.
;                   MORENO VALLE,Migeul Angel
;                   SALTO MALDONADO, Francisco
;                   DIEZ GARCIA, Bruno
;        TITLE OF INVENTION: PROMOTERS OF THE GENES GLUTAMATE
;                            DESHYDROGENASE,  -N-ACETYLHEXOSAMINIDASE
;                            AND  -ACTIN AND THEIR USE IN FILAMENTOUS
;                            FUNGI EXPRESSION, SECRETION AND ANTISENSE
;        NUMBER OF SEQUENCES: 20
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: LADAS & PARRY
;             STREET: 26 WEST 61 STREET
;             CITY: NEW YORK
;             STATE: NY
;             COUNTRY: USA
;             ZIP: 10023
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: 3-1/4" Disk  1.44MB
;
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```
;             COMPUTER: IBM PC compatible
;             OPERATING SYSTEM: Microsoft Windows for Workgroups 3.11
;             SOFTWARE: WordPerfect 8 for Windows
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/171,337A
;             FILING DATE: 14-May-1999
;             CLASSIFICATION: 536
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: PCT/ES98/00056
;             FILING DATE: 5-MAR-1998
;             APPLICATION NUMBER: ES9700482
;             FILING DATE: 5-MAR-1997
;        ATTORNEY/AGENT INFORMATION:
;             NAME: MASS, Clifford J.
;             REGISTRATION NUMBER: 30,086
;                                  (C) REF./DOCKET NO.:    U-011948-3
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: 233288
;   INFORMATION FOR SEQ ID NO: 6
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 596 amino acids
;             TYPE: amino acids
;             STRANDEDNESS: single
;             TOPOLOGY: linear
;        MOLECULE TYPE: peptide
;        ORIGINAL SOURCE:
;             ORGANISM: Penicillum chrysogenum
;        FEATURE:
;             OTHER INFORMATION:    amino acid sequence of the  -N-
;             acetylhexosaminidase enzyme
;             (EC.3.2.1.52) with a
;             molecular weight of 66545 Da.
;        SEQUENCE DESCRIPTION: SEQ ID NO: 6
US-09-171-337A-6

  Query Match             50.7%;  Score 37.5;  DB 2;  Length 596;
  Best Local Similarity   81.8%;  Pred. No. 1.2e+02;
  Matches    9;  Conservative    1;  Mismatches    0;  Indels    1;  Gaps    1;

Qy          4 GMALSKIN-LH 13
              ||||||:| ||
Db        202 GMALSKLNVLH 212


RESULT 32
US-09-631-022-6
; Sequence 6, Application US/09631022
; Patent No. 6558921
;   GENERAL INFORMATION:
;        APPLICANT: BARREDO FUENTE, Jose Luis
;                   RODRIGUEZ SAIZ, Marta
;                   COLLADOS DE LA VIEJA, Alfonso J.
;                   MORENO VALLE,Migeul Angel
;                   SALTO MALDONADO, Francisco
;                   DIEZ GARCIA, Bruno
;        TITLE OF INVENTION: PROMOTERS OF THE GENES GLUTAMATE
;                            DESHYDROGENASE,  -N-ACETYLHEXOSAMINIDASE
;                            AND  -ACTIN AND THEIR USE IN FILAMENTOUS
;                            FUNGI EXPRESSION, SECRETION AND ANTISENSE
;        NUMBER OF SEQUENCES: 20
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: LADAS & PARRY
;             STREET: 26 WEST 61 STREET
;             CITY: NEW YORK
;             STATE: NY
;             COUNTRY: USA
;             ZIP: 10023
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: 3-1/4" Disk  1.44MB
;             COMPUTER: IBM PC compatible
;             OPERATING SYSTEM: Microsoft Windows for Workgroups 3.11
;             SOFTWARE: WordPerfect 8 for Windows
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/631,022
;             FILING DATE: 02-Aug-2000
;             CLASSIFICATION: 536
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: 09/171,337
;             FILING DATE: 14-MAY-1999
;             APPLICATION NUMBER: PCT/ES98/00056
;             FILING DATE: 5-MAR-1998
;             APPLICATION NUMBER: ES9700482
;             FILING DATE: 5-MAR-1997
;        ATTORNEY/AGENT INFORMATION:
;             NAME: MASS, Clifford J.
;             REGISTRATION NUMBER: 30,086
;                                  (C) REF./DOCKET NO.:    U-02886-6
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: 233288
;   INFORMATION FOR SEQ ID NO: 6
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 596 amino acids
;             TYPE: amino acids
;             STRANDEDNESS: single
;             TOPOLOGY: linear
;        MOLECULE TYPE: peptide
;        ORIGINAL SOURCE:
;             ORGANISM: Penicillum chrysogenum
;        FEATURE:
;             OTHER INFORMATION:    amino acid sequence of the  -N-
;             acetylhexosaminidase enzyme
;             (EC.3.2.1.52) with a
;             molecular weight of 66545 Da.
;        SEQUENCE DESCRIPTION: SEQ ID NO: 6
US-09-631-022-6

  Query Match             50.7%;  Score 37.5;  DB 2;  Length 596;
  Best Local Similarity   81.8%;  Pred. No. 1.2e+02;
  Matches    9;  Conservative    1;  Mismatches    0;  Indels    1;  Gaps    1;

Qy          4 GMALSKIN-LH 13
              ||||||:| ||
Db        202 GMALSKLNVLH 212


RESULT 33
US-09-830-807-7
; Sequence 7, Application US/09830807
; Patent No. 6846667
; GENERAL INFORMATION:
;  APPLICANT: Crooke,Helen R.
;  APPLICANT:  Clarke, Enda E.
;  APPLICANT:  Everest, Paul H.
;  APPLICANT:  Dougan, Gordon
;  APPLICANT:  Holden, David W.
;  APPLICANT:  Shea, Jacqueline E.
;  APPLICANT:  Feldman, Robert G.
;  TITLE OF INVENTION: VIRULENCE GENES AND PROTEINS, AND THEIR USE
;  FILE REFERENCE: GJE-65
;  CURRENT APPLICATION NUMBER: US/09/830,807
;  CURRENT FILING DATE:  2001-04-30
;  NUMBER OF SEQ ID NOS: 72
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 7
;   LENGTH: 42
;   TYPE: PRT
;   ORGANISM: Escherichia coli
US-09-830-807-7

  Query Match             50.0%;  Score 37;  DB 2;  Length 42;
  Best Local Similarity   62.5%;  Pred. No. 7.2;
  Matches   10;  Conservative    0;  Mismatches    2;  Indels    4;  Gaps    1;

Qy          2 GYGMALS----KINLH 13
              | | |||    |||||
Db         16 GVGAALSNKLAKINLH 31
```

```
RESULT 34
US-09-270-767-37334
; Sequence 37334, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
;  APPLICANT: Homburger et al.
;  TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
;  FILE REFERENCE: File Reference: 7326-094
;  CURRENT APPLICATION NUMBER: US/09/270,767
;  CURRENT FILING DATE: 1999-03-17
;  NUMBER OF SEQ ID NOS: 62517
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 37334
;   LENGTH: 167
;   TYPE: PRT
;   ORGANISM: Drosophila melanogaster
US-09-270-767-37334

  Query Match             50.0%;  Score 37;  DB 2;  Length 167;
  Best Local Similarity   61.5%;  Pred. No. 35;
  Matches    8;  Conservative    1;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              |||  || ||: |
Db         66 GYGDILSGINIAN 78


RESULT 35
US-09-270-767-52551
; Sequence 52551, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
;  APPLICANT: Homburger et al.
;  TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
;  FILE REFERENCE: File Reference: 7326-094
;  CURRENT APPLICATION NUMBER: US/09/270,767
;  CURRENT FILING DATE: 1999-03-17
;  NUMBER OF SEQ ID NOS: 62517
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 52551
;   LENGTH: 167
;   TYPE: PRT
;   ORGANISM: Drosophila melanogaster
US-09-270-767-52551

  Query Match             50.0%;  Score 37;  DB 2;  Length 167;
  Best Local Similarity   61.5%;  Pred. No. 35;
  Matches    8;  Conservative    1;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              |||  || ||: |
Db         66 GYGDILSGINIAN 78


RESULT 36
US-09-949-016-6509
; Sequence 6509, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 6509
;   LENGTH: 217
;   TYPE: PRT
;   ORGANISM: Human
US-09-949-016-6509

  Query Match             50.0%;  Score 37;  DB 2;  Length 217;
  Best Local Similarity   50.0%;  Pred. No. 47;
  Matches    6;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLH 13
              |||:  |  |:|
Db         77 GYGLEPSSFNMH 88


RESULT 37
US-09-949-016-9719
; Sequence 9719, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 9719
;   LENGTH: 228
;   TYPE: PRT
;   ORGANISM: Human
US-09-949-016-9719

  Query Match             50.0%;  Score 37;  DB 2;  Length 228;
  Best Local Similarity   50.0%;  Pred. No. 49;
  Matches    6;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLH 13
              |||:  |  |:|
Db         88 GYGLEPSSFNMH 99


RESULT 38
US-09-134-000C-6538
; Sequence 6538, Application US/09134000C
; Patent No. 6617156
; GENERAL INFORMATION:
;  APPLICANT: Lynn Doucette-Stamm et al
;  TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
;  TITLE OF INVENTION:  ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS
;  FILE REFERENCE: 032796-032
;  CURRENT APPLICATION NUMBER: US/09/134,000C
;  CURRENT FILING DATE:  1998-08-13
;  PRIOR APPLICATION NUMBER: US 60/055,778
;  PRIOR FILING DATE: 1997-08-15
;  NUMBER OF SEQ ID NOS: 6812
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 6538
;   LENGTH: 618
;   TYPE: PRT
;   ORGANISM: Enterococcus faecalis
US-09-134-000C-6538

  Query Match             50.0%;  Score 37;  DB 2;  Length 618;
```

```
  Best Local Similarity   53.8%;  Pred. No. 1.5e+02;
  Matches    7;  Conservative    1;  Mismatches    5;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              |||   |  :|| |
Db        389 GYGYMLYSVNLKN 401


RESULT 39
US-09-949-016-6354
; Sequence 6354, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 6354
;   LENGTH: 351
;   TYPE: PRT
;   ORGANISM: Human
US-09-949-016-6354

  Query Match             49.3%;  Score 36.5;  DB 2;  Length 351;
  Best Local Similarity   31.0%;  Pred. No. 1e+02;
  Matches    9;  Conservative    4;  Mismatches    1;  Indels   15;  Gaps    1;

Qy          1 MGYGMALSK--------------INLHN 14
              :|:| |:||               :||||
Db        136 VGFGEAISKQFVDALETGQDARAAMNLHN 164


RESULT 40
US-09-949-016-7724
; Sequence 7724, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
;  APPLICANT: VENTER, J. Craig et al.
;  TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
;  TITLE OF INVENTION:  WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001307
;  CURRENT APPLICATION NUMBER: US/09/949,016
;  CURRENT FILING DATE:  2000-04-14
;  PRIOR APPLICATION NUMBER: 60/241,755
;  PRIOR FILING DATE: 2000-10-20
;  PRIOR APPLICATION NUMBER: 60/237,768
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/231,498
;  PRIOR FILING DATE: 2000-09-08
;  NUMBER OF SEQ ID NOS: 207012
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 7724
;   LENGTH: 357
;   TYPE: PRT
;   ORGANISM: Human
US-09-949-016-7724

  Query Match             49.3%;  Score 36.5;  DB 2;  Length 357;
  Best Local Similarity   31.0%;  Pred. No. 1e+02;
  Matches    9;  Conservative    4;  Mismatches    1;  Indels   15;  Gaps    1;

Qy          1 MGYGMALSK--------------INLHN 14
              :|:| |:||               :||||
Db        142 VGFGEAISKQFVDALETGQDARAAMNLHN 170


Search completed: May 13, 2006, 08:15:31
Job time : 48 secs
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GenCore version 5.1.8

OM protein - protein search, using sw model

Run on: May 13, 2006, 08:06:48 ; Search time 185 Seconds
(without alignments)
33.250 Million cell updates/sec

Title: US-10-769-514-17
Perfect score: 74
Sequence: 1 MGYGMALSKINLHN 14

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2443163 seqs, 439378781 residues

Total number of hits satisfying chosen parameters: 2443163

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 100 summaries

Database : A_Geneseq_21:*
1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
5: geneseqp2002s:*
6: geneseqp2003as:*
7: geneseqp2003bs:*
8: geneseqp2004s:*
9: geneseqp2005s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	74	100.0	14	9	ADV70480	Adv70480 Moraxella
2	74	100.0	15	9	ADV70478	Adv70478 Moraxella
3	74	100.0	15	9	ADV70479	Adv70479 Moraxella
4	74	100.0	702	2	AAW35313	Aaw35313 M. catarr
5	74	100.0	702	2	AAY43383	Aay43383 M. catarr
6	65	87.8	706	2	AAY43378	Aay43378 M. catarr
7	63	85.1	714	2	AAW35316	Aaw35316 M. catarr
8	46	62.2	190	6	ABM69725	Abm69725 Photorhab
9	42	56.8	635	8	ADI36920	Adi36920 Human LRR
10	42	56.8	695	3	AAY94963	Aay94963 Human sec
11	42	56.8	696	2	AAY08076	Aay08076 Human PRO
12	42	56.8	696	2	AAY13359	Aay13359 Amino aci
13	42	56.8	696	3	AAY70671	Aay70671 Human PRO
14	42	56.8	696	3	ADC78411	Adc78411 Human PRO
15	42	56.8	696	4	AAB80227	Aab80227 Human PRO
16	42	56.8	696	4	AAU00824	Aau00824 Human imm
17	42	56.8	696	4	AAU12348	Aau12348 Human PRO
18	42	56.8	696	4	AAB50905	Aab50905 Human PRO
19	42	56.8	696	6	ABU71605	Abu71605 Human PRO
20	42	56.8	696	6	ABO17792	Abo17792 Novel hum
21	42	56.8	696	6	ABU71460	Abu71460 Human PRO
22	42	56.8	696	6	ABU81046	Abu81046 Human PRO
23	42	56.8	696	6	ABU71906	Abu71906 Human sec
24	42	56.8	696	6	ABO01789	Abo01789 Novel hum
25	42	56.8	696	6	ABU66746	Abu66746 Human PRO
26	42	56.8	696	6	ABU54362	Abu54362 Human sec
27	42	56.8	696	6	ABO47377	Abo47377 Human sec
28	42	56.8	696	6	ABU59827	Abu59827 Novel sec
29	42	56.8	696	6	ABO25017	Abo25017 Human sec
30	42	56.8	696	6	ABU64514	Abu64514 Human sec
31	42	56.8	696	6	ABU67360	Abu67360 Human sec
32	42	56.8	696	6	ABO14880	Abo14880 Human sec
33	42	56.8	696	6	ABU67022	Abu67022 Human sec
34	42	56.8	696	6	ABU69637	Abu69637 Novel hum
35	42	56.8	696	6	ABO14819	Abo14819 Human sec
36	42	56.8	696	6	ADA45873	Ada45873 Novel hum
37	42	56.8	696	6	ADA76304	Ada76304 Human PRO
38	42	56.8	696	6	ADB29296	Adb29296 Human sec
39	42	56.8	696	6	ADA18954	Ada18954 Human PRO
40	42	56.8	696	6	ADA61577	Ada61577 Homo sapi
41	42	56.8	696	6	ADB19362	Adb19362 Novel hum
42	42	56.8	696	6	ADB27903	Adb27903 Human PRO
43	42	56.8	696	6	ADA86382	Ada86382 Novel hum
44	42	56.8	696	6	ADB15946	Adb15946 Human PRO
45	42	56.8	696	6	ADA47732	Ada47732 Human PRO
46	42	56.8	696	6	ADA18152	Ada18152 Human sec
47	42	56.8	696	6	ABO32771	Abo32771 Human sec
48	42	56.8	696	6	ADA67527	Ada67527 Human PRO
49	42	56.8	696	6	ADB30534	Adb30534 Human PRO
50	42	56.8	696	6	ADA85830	Ada85830 Novel hum
51	42	56.8	696	6	ADA97042	Ada97042 Human PRO
52	42	56.8	696	6	ADA79346	Ada79346 Human PRO
53	42	56.8	696	6	ADA87485	Ada87485 Novel hum
54	42	56.8	696	6	ADB16687	Adb16687 Human PRO
55	42	56.8	696	6	ABO34831	Abo34831 Human PRO
56	42	56.8	696	6	ADA16127	Ada16127 Human sec
57	42	56.8	696	6	ADA91779	Ada91779 Novel hum
58	42	56.8	696	6	ADB14842	Adb14842 Human PRO
59	42	56.8	696	6	ADB18803	Adb18803 Novel hum
60	42	56.8	696	6	ADA94018	Ada94018 Human PRO
61	42	56.8	696	6	ADB19914	Adb19914 Novel hum
62	42	56.8	696	6	ADB13226	Adb13226 Human PRO
63	42	56.8	696	6	ABO43325	Abo43325 Novel hum
64	42	56.8	696	6	ADA74480	Ada74480 Human PRO
65	42	56.8	696	6	ADA42272	Ada42272 Human sec
66	42	56.8	696	6	ADB24713	Adb24713 Human PRO
67	42	56.8	696	6	ADA82237	Ada82237 Human PRO
68	42	56.8	696	6	ADA75200	Ada75200 Human PRO
69	42	56.8	696	6	ADA85278	Ada85278 Novel hum
70	42	56.8	696	6	ADA84726	Ada84726 Novel hum
71	42	56.8	696	6	ABO17509	Abo17509 Human PRO
72	42	56.8	696	6	ADB29982	Adb29982 Human PRO
73	42	56.8	696	6	ADA80510	Ada80510 Human PRO
74	42	56.8	696	6	ADA75752	Ada75752 Human PRO
75	42	56.8	696	6	ADA46977	Ada46977 Human PRO
76	42	56.8	696	6	ADB25273	Adb25273 Human PRO
77	42	56.8	696	6	ADA93449	Ada93449 Human PRO
78	42	56.8	696	6	ADB26799	Adb26799 Human PRO
79	42	56.8	696	6	ADB31086	Adb31086 Human PRO
80	42	56.8	696	6	ADA61014	Ada61014 Homo sapi
81	42	56.8	696	6	ADB24161	Adb24161 Human PRO
82	42	56.8	696	6	ADA96490	Ada96490 Human PRO
83	42	56.8	696	6	ADA81062	Ada81062 Human PRO
84	42	56.8	696	6	ADA95938	Ada95938 Human PRO
85	42	56.8	696	6	ADB26247	Adb26247 Human PRO
86	42	56.8	696	6	ADB21732	Adb21732 Novel hum
87	42	56.8	696	7	ADA77511	Ada77511 Human PRO
88	42	56.8	696	7	ADB18251	Adb18251 Human PRO
89	42	56.8	696	7	ADA86934	Ada86934 Novel hum
90	42	56.8	696	7	ADA16551	Ada16551 Human sec
91	42	56.8	696	7	ADA12980	Ada12980 Human sec
92	42	56.8	696	7	ADA41848	Ada41848 Human sec
93	42	56.8	696	7	ADA88037	Ada88037 Novel hum
94	42	56.8	696	7	ADA46425	Ada46425 Novel hum
95	42	56.8	696	7	ADA17195	Ada17195 Human sec
96	42	56.8	696	7	ADA42698	Ada42698 Human sec
97	42	56.8	696	7	ADB28455	Adb28455 Human PRO

98	42	56.8	696	7	ADB29007	Adb29007 Human PRO
99	42	56.8	696	7	ADA76959	Ada76959 Human PRO
100	42	56.8	696	7	ADA88589	Ada88589 Novel hum

ALIGNMENTS

RESULT 1
ADV70480
ID ADV70480 standard; peptide; 14 AA.
XX
AC ADV70480;
XX
DT 10-MAR-2005 (first entry)
XX
DE Moraxella catarrhalis transferrin binding protein B region - SEQ ID 17.
XX
KW bacterial infection; antibacterial; bacterial meningitis; antibacterial;
KW neuroprotective; otitis media; auditory; transferrin binding protein B.
XX
OS Moraxella catarrhalis.
XX
PN US2004258695-A1.
XX
PD 23-DEC-2004.
XX
PF 30-JAN-2004; 2004US-00769514.
XX
PR 31-JAN-2003; 2003US-0444113P.
XX
PA (SCHR/) SCHRYVERS A B.
XX
PI Schryvers AB;
XX
DR WPI; 2005-038740/04.
XX
PT Transferrin-binding molecules useful for eliciting antibodies to
PT bacterial transferrin binding proteins, which block bacterial transferrin
PT uptake.
XX
PS Claim 7; SEQ ID NO 17; 27pp; English.
XX
CC The invention comprises a molecule (e.g. peptide) which is capable of:
CC binding to a region of a transferrin protein that is recognized by a
CC bacterial transferrin binding protein; and eliciting an antibody to the
CC bacterial transferrin binding protein. The transferrin binding molecule
CC of the invention is useful for preventing and treating bacterial
CC infections (e.g. bacterial meningitis and otitis media). The present
CC amino acid sequence represents a region of the Moraxella catarrhalis
CC transferrin binding protein B.
XX
SQ Sequence 14 AA;

Query Match 100.0%; Score 74; DB 9; Length 14;
Best Local Similarity 100.0%; Pred. No. 5.9e-07;
Matches 14; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
||||||||||||||
Db 1 MGYGMALSKINLHN 14

RESULT 2
ADV70478
ID ADV70478 standard; peptide; 15 AA.
XX
AC ADV70478;
XX
DT 10-MAR-2005 (first entry)
XX
DE Moraxella catarrhalis transferrin binding protein B region - SEQ ID 15.
XX
KW bacterial infection; antibacterial; bacterial meningitis; antibacterial;
KW neuroprotective; otitis media; auditory; transferrin binding protein B.
XX
OS Moraxella catarrhalis.
XX
PN US2004258695-A1.
XX
PD 23-DEC-2004.
XX
PF 30-JAN-2004; 2004US-00769514.
XX
PR 31-JAN-2003; 2003US-0444113P.
XX
PA (SCHR/) SCHRYVERS A B.
XX
PI Schryvers AB;
XX
DR WPI; 2005-038740/04.
XX
PT Transferrin-binding molecules useful for eliciting antibodies to
PT bacterial transferrin binding proteins, which block bacterial transferrin
PT uptake.
XX
PS Example 1; SEQ ID NO 15; 27pp; English.
XX
CC The invention comprises a molecule (e.g. peptide) which is capable of:
CC binding to a region of a transferrin protein that is recognized by a
CC bacterial transferrin binding protein; and eliciting an antibody to the
CC bacterial transferrin binding protein. The transferrin binding molecule
CC of the invention is useful for preventing and treating bacterial
CC infections (e.g. bacterial meningitis and otitis media). The present
CC amino acid sequence represents a region of the Moraxella catarrhalis
CC transferrin binding protein B.
XX
SQ Sequence 15 AA;

Query Match 100.0%; Score 74; DB 9; Length 15;
Best Local Similarity 100.0%; Pred. No. 6.4e-07;
Matches 14; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
||||||||||||||
Db 1 MGYGMALSKINLHN 14

RESULT 3
ADV70479
ID ADV70479 standard; peptide; 15 AA.
XX
AC ADV70479;
XX
DT 10-MAR-2005 (first entry)
XX
DE Moraxella catarrhalis transferrin binding protein B region - SEQ ID 16.
XX
KW bacterial infection; antibacterial; bacterial meningitis; antibacterial;
KW neuroprotective; otitis media; auditory; transferrin binding protein B.
XX
OS Moraxella catarrhalis.
XX
PN US2004258695-A1.
XX
PD 23-DEC-2004.
XX
PF 30-JAN-2004; 2004US-00769514.
XX
PR 31-JAN-2003; 2003US-0444113P.
XX
PA (SCHR/) SCHRYVERS A B.
XX
PI Schryvers AB;
XX
DR WPI; 2005-038740/04.

XX
PT Transferrin-binding molecules useful for eliciting antibodies to
PT bacterial transferrin binding proteins, which block bacterial transferrin
PT uptake.
XX
PS Example 1; SEQ ID NO 16; 27pp; English.
XX
CC The invention comprises a molecule (e.g. peptide) which is capable of:
CC binding to a region of a transferrin protein that is recognized by a
CC bacterial transferrin binding protein; and eliciting an antibody to the
CC bacterial transferrin binding protein. The transferrin binding molecule
CC of the invention is useful for preventing and treating bacterial
CC infections (e.g. bacterial meningitis and otitis media). The present
CC amino acid sequence represents a region of the Moraxella catarrhalis
CC transferrin binding protein B.
XX
SQ Sequence 15 AA;

Query Match 100.0%; Score 74; DB 9; Length 15;
Best Local Similarity 100.0%; Pred. No. 6.4e-07;
Matches 14; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
||||||||||||||
Db 2 MGYGMALSKINLHN 15

RESULT 4
AAW35313
ID AAW35313 standard; protein; 702 AA.
XX
AC AAW35313;
XX
DT 14-APR-1998 (first entry)
XX
DE M. catarrhalis 4223 transferrin binding protein tbpB.
XX
KW Transferrin binding protein; tbpB; immunogen; vaccine; protection;
KW otitis media; antibody; diagnosis; therapy; carrier; gene isolation.
XX
OS Moraxella catarrhalis.
XX
PN WO9732980-A1.
XX
PD 12-SEP-1997.
XX
PF 07-MAR-1997; 97WO-CA000163.
XX
PR 08-MAR-1996; 96US-00613009.
PR 03-JAN-1997; 97US-00778570.
XX
PA (CONN-) CONNAUGHT LAB LTD.
XX
PI Myers LE, Schryvers AB, Harkness RE, Loosmore SM, Du R, Yang Y;
PI Klein MH;
XX
DR WPI; 1997-457533/42.
DR N-PSDB; AAT95248.
XX
PT DNA encoding transferrin receptor of a Moraxella strain - also proteins,
PT useful in vaccines, as diagnostic agents and in the production of
PT antibodies.
XX
PS Claim 6; Fig 6; 162pp; English.
XX
CC The present sequence is the Moraxella catarrhalis 4223 transferrin
CC binding protein tbpB, which can be used as an immunogen, e.g. in vaccines
CC to protect against diseases caused by M. catarrhalis (specifically otitis
CC media), or to raise antibodies for diagnosis and therapy. It can also be
CC used as a carrier for other antigenic determinants, e.g. of bacteria
CC containing polysaccharide antigens or abnormal polysaccharides present on
CC tumour cells, particularly to make conjugate vaccines. The tbpB DNA can
CC be used to detect nucleic acid encoding transferrin receptor protein,
CC e.g. for diagnosis or gene isolation, by usual hybridisation assays
XX
SQ Sequence 702 AA;

Query Match 100.0%; Score 74; DB 2; Length 702;
Best Local Similarity 100.0%; Pred. No. 5.2e-05;
Matches 14; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
||||||||||||||
Db 100 MGYGMALSKINLHN 113

RESULT 5
AAY43383
ID AAY43383 standard; protein; 702 AA.
XX
AC AAY43383;
XX
DT 26-JAN-2000 (first entry)
XX
DE M. catarrhalis strain 4223 tbp2 protein.
XX
KW TbpB gene; Tbp2; transferrin binding protein; diagnosis; otitis media;
KW genetic immunisation; Moraxella infection; antigen; vaccine; detection;
KW antitumour antibody production; therapy.
XX
OS Moraxella catarrhalis.
XX
PN WO9952947-A2.
XX
PD 21-OCT-1999.
XX
PF 12-APR-1999; 99WO-CA000307.
XX
PR 14-APR-1998; 98US-00059584.
XX
PA (CONN-) CONNAUGHT LAB LTD.
XX
PI Myers LE, Schryvers AB, Harkness RE, Loosmore SM, Du R, Yang Y;
PI Klein MH;
XX
DR WPI; 1999-620376/53.
DR N-PSDB; AAZ31949.
XX
PT Nucleic acid encoding transferrin binding protein 2 of Moraxella
PT catarrhalis, useful for diagnostics, immunization and recombinant protein
PT production.
XX
PS Example 6; Fig 8; 114pp; English.
XX
CC This sequence is the Moraxella catarrhalis strain 4223 transferrin
CC binding protein (Tbp2) of the invention. The DNA sequence is also
CC referred to as the TbpB gene. The TbpB gene is used to produce
CC recombinant Tbp2; for identification or diagnosis of Moraxella, or for
CC cloning related species, using hybridisation assays; and for genetic
CC immunisation against Moraxella infections, e.g. otitis media. The Tbp2
CC proteins are useful as antigens, either in vaccines (including components
CC of conjugate vaccines that contain antigens from other bacteria or from
CC tumours, in which case they elicit production of antitumour antibodies
CC that may be coupled to chemotherapeutic agents or biologically active
CC agents) or to raise antibodies (for use as diagnostic reagents and for
CC treating Moraxella infections), also for detecting Moraxella antibodies
XX
SQ Sequence 702 AA;

Query Match 100.0%; Score 74; DB 2; Length 702;
Best Local Similarity 100.0%; Pred. No. 5.2e-05;
Matches 14; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
||||||||||||||
Db 100 MGYGMALSKINLHN 113

```
RESULT 6
AAY43378
ID   AAY43378 standard; protein; 706 AA.
XX
AC   AAY43378;
XX
DT   26-JAN-2000  (first entry)
XX
DE   M. catarrhalis strain M35 tbp2 protein.
XX
KW   TbpB gene; Tbp2; transferrin binding protein; diagnosis; otitis media;
KW   genetic immunisation; Moraxella infection; antigen; vaccine; detection;
KW   antitumour antibody production; therapy.
XX
OS   Moraxella catarrhalis.
XX
PN   WO9952947-A2.
XX
PD   21-OCT-1999.
XX
PF   12-APR-1999;   99WO-CA000307.
XX
PR   14-APR-1998;   98US-00059584.
XX
PA   (CONN-) CONNAUGHT LAB LTD.
XX
PI   Myers LE,  Schryvers AB,  Harkness RE,  Loosmore SM,  Du R,  Yang Y;
PI   Klein MH;
XX
DR   WPI; 1999-620376/53.
DR   N-PSDB; AAZ31946.
XX
PT   Nucleic acid encoding transferrin binding protein 2 of Moraxella
PT   catarrhalis, useful for diagnostics, immunization and recombinant protein
PT   production.
XX
PS   Claim 7; Fig 2; 114pp; English.
XX
CC   This sequence is the Moraxella catarrhalis strain M35 transferrin binding
CC   protein (Tbp2) of the invention. The DNA sequence is also referred to as
CC   the TbpB gene. The TbpB gene is used to produce recombinant Tbp2; for
CC   identification or diagnosis of Moraxella, or for cloning related species,
CC   using hybridisation assays; and for genetic immunisation against
CC   Moraxella infections, e.g. otitis media. The Tbp2 proteins are useful as
CC   antigens, either in vaccines (including components of conjugate vaccines
CC   that contain antigens from other bacteria or from tumours, in which case
CC   they elicit production of antitumour antibodies that may be coupled to
CC   chemotherapeutic agents or biologically active agents) or to raise
CC   antibodies (for use as diagnostic reagents and for treating Moraxella
CC   infections), also for detecting Moraxella antibodies
XX
SQ   Sequence 706 AA;

  Query Match             87.8%;  Score 65;  DB 2;  Length 706;
  Best Local Similarity   92.3%;  Pred. No. 0.0028;
  Matches   12;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLH 13
              |||||||||:|||
Db        100 MGYGMALSKMNLH 112


RESULT 7
AAW35316
ID   AAW35316 standard; protein; 714 AA.
XX
AC   AAW35316;
XX
DT   14-APR-1998  (first entry)
XX
DE   M. catarrhalis R1 transferrin binding protein tbpB.
XX
KW   Transferrin binding protein; tbpB; immunogen; vaccine; protection;
KW   otitis media; antibody; diagnosis; therapy; carrier; gene isolation.
XX
OS   Moraxella catarrhalis.
XX
PN   WO9732980-A1.
XX
PD   12-SEP-1997.
XX
PF   07-MAR-1997;   97WO-CA000163.
XX
PR   08-MAR-1996;   96US-00613009.
PR   03-JAN-1997;   97US-00778570.
XX
PA   (CONN-) CONNAUGHT LAB LTD.
XX
PI   Myers LE,  Schryvers AB,  Harkness RE,  Loosmore SM,  Du R,  Yang Y;
PI   Klein MH;
XX
DR   WPI; 1997-457533/42.
DR   N-PSDB; AAT95251.
XX
PT   DNA encoding transferrin receptor of a Moraxella strain - also proteins,
PT   useful in vaccines, as diagnostic agents and in the production of
PT   antibodies.
XX
PS   Claim 6; Fig 27; 162pp; English.
XX
CC   The present sequence is the Moraxella catarrhalis R1 transferrin binding
CC   protein tbpB, which can be used as an immunogen, e.g. in vaccines to
CC   protect against diseases caused by M. catarrhalis (specifically otitis
CC   media), or to raise antibodies for diagnosis and therapy. It can also be
CC   used as a carrier for other antigenic determinants, e.g. of bacteria
CC   containing polysaccharide antigens or abnormal polysaccharides present on
CC   tumour cells, particularly to make conjugate vaccines. The tbpB DNA can
CC   be used to detect nucleic acid encoding transferrin receptor protein,
CC   e.g. for diagnosis or gene isolation, by usual hybridisation assays
XX
SQ   Sequence 714 AA;

  Query Match             85.1%;  Score 63;  DB 2;  Length 714;
  Best Local Similarity   85.7%;  Pred. No. 0.0067;
  Matches   12;  Conservative    2;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||::
Db        100 MGYGMALSKINLYD 113


RESULT 8
ABM69725
ID   ABM69725 standard; protein; 190 AA.
XX
AC   ABM69725;
XX
DT   20-NOV-2003  (first entry)
XX
DE   Photorhabdus luminescens protein sequence #2822.
XX
KW   Antibacterial; fungicide; insecticide; polymorphism; genetic analysis;
KW   detection; food; gene expression; plant; animal; microorganism; toxin;
KW   antibiotic; biopesticide; virulence factor; disease model; plague;
KW   whooping cough.
XX
OS   Photorhabdus luminescens.
XX
PN   WO200294867-A2.
XX
PD   28-NOV-2002.
XX
PF   07-FEB-2002; 2002WO-IB003040.
XX
```

PR 07-FEB-2001; 2001FR-00001659.
XX
PA (INSP) INST PASTEUR.
PA (CNRS) CNRS CENT NAT RECH SCI.
XX
PI Duchaud E, Taourit S, Glaser P, Frangeul L, Kunst F, Danchin A;
PI Buchrieser C;
XX
DR WPI; 2003-148459/14.
XX
PT Genomic sequence of Photorhabdus luminescens and encoded polypeptides,
PT useful e.g. as therapeutic antimicrobials and agricultural pesticides.
XX
PS Claim 2; SEQ ID NO 2822; 1205pp; French.
XX
CC The invention relates to the isolation of genes and their encoded
CC proteins from Photorhabdus luminescens. The isolated sequences are
CC sources of probes and primers for detecting the genome of P. luminescens
CC and related species; to study polymorphisms; for gene analysis and for
CC detection/amplification of the genes. Antibodies (Ab) raised against the
CC polypeptides encoded by the genes are used for detection/identification
CC of P. luminescens, e.g. in foods. The genes, proteins, Ab and cells that
CC carry a gene-containing vector are used to select compounds that
CC modulate, regulate, induce or inhibit expression of the genes in plants,
CC animals or microorganisms other than P. luminescens and are able to alter
CC response or sensitivity to toxins and antibiotics produced by P.
CC luminescens. Cells transformed to express the genes are useful for
CC recombinant production of the proteins, particularly toxins and
CC antibacterials useful as insecticides, bactericides and fungicides. The
CC genes, proteins, vectors containing the genes and Ab are also useful
CC therapeutically (to treat microbial infection by bacteria or fungi that
CC are sensitive to P. luminescens-encoded toxins or antibiotics) and as
CC biopesticides. Other uses of the genes and the proteins are as virulence
CC factors and for identifying targets of human diseases for which P.
CC luminescens is a model (particularly plague and whooping cough). This
CC sequence represents one of the isolated P. luminescens proteins
XX
SQ Sequence 190 AA;

Query Match 62.2%; Score 46; DB 6; Length 190;
Best Local Similarity 57.1%; Pred. No. 2.6;
Matches 8; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
||:|| :||||
Db 15 MGFGMTTILLNLHN 28

RESULT 9
ADI36920
ID ADI36920 standard; protein; 635 AA.
XX
AC ADI36920;
XX
DT 22-APR-2004 (first entry)
XX
DE Human LRR protein #15.
XX
KW Leucine-rich repeat; LRR; HLRRNS-2; HLRRNS-3;
KW aberrant leucine-rich repeat protein function disorder;
KW protein: protein interaction disorder; matrix association disorder;
KW caspase recruitment disorder; nucleotide binding disorder;
KW cell migration disorder; signal transduction disorder;
KW cell cycle regulation disorder; neurological disorder;
KW motor neuron disorder; muscle development disorder;
KW neural development disorder; apoptosis disorder;
KW immune response disorder; dementia; anxiety; headache; migraine;
KW delirium; schizophrenia; manic depression; mental retardation;
KW dyskinesia; neural degenerative disorder; Alzheimers's disease;
KW Parkinson's disease; depression; fear; learning disorder; brain cancer;
KW gene therapy; human; nootropic; tranquilizer; neuroleptic;
KW neuroprotective; cytostatic.
XX
OS Homo sapiens.
XX
PN US2003220263-A1.
XX
PD 27-NOV-2003.
XX
PF 25-APR-2003; 2003US-00424233.
XX
PR 25-APR-2002; 2002US-0375335P.
XX
PA (FEDE/) FEDER J N.
PA (MINT/) MINTIER G.
PA (RAMA/) RAMANATHAN C S.
XX
PI Feder JN, Mintier G, Ramanathan CS;
XX
DR WPI; 2004-141759/14.
XX
PT New isolated human leucine rich repeat containing polypeptides, HLRRNS-2
PT and HLRRNS-3, useful for treating, preventing disorders e.g., anxiety,
PT headache, migraine, schizophrenia, manic depression, or delirium.
XX
PS Example 1; SEQ ID NO 19; 124pp; English.
XX
CC The present invention relates to two newly described human leucine-rich
CC repeat (LRR) containing proteins HLRRNS-2 and HLRRNS-3 and their encoding
CC nucleic acids. The invention is useful for preventing, treating and
CC ameliorating a medical condition which involves administeration of LRR
CC proteins or their modulators . The invention is also useful for
CC diagnosing a pathological condition or a susceptibility of the
CC pathological condition which involves determining the presence or amount
CC of expression of the LRR proteins in a biological sample. The condition
CC is disorder related to aberrant leucine-rich repeat protein function, a
CC disorder related to aberrant protein: protein interactions, disorders
CC related to aberrant matrix association, a disorder related to aberrant
CC caspase recruitment, a disorder related to aberrant nucleotide binding, a
CC disorder related to aberrant cell migration, a disorder related to
CC aberrant signal transduction, a disorder related to aberrant cell cycle
CC regulation, neurological disorders, motor neuron disorders, muscle
CC development disorders, a disorder related to aberrant neural development,
CC a disorder related to aberrant apoptosis, disorder related to aberrant
CC immune responses in the human nervous system such as dementia, anxiety,
CC headache, migraine, delirium, schizophrenia, manic depression, severe
CC mental retardation, dyskinesias, neural degenerative disorders such as
CC Alzheimers's disease, Parkinson's disease affective disorders,
CC depression, schizophrenia, anxiety, fear, learning disorders and brain
CC cancer. The invention is also used in gene therapy. The present sequence
CC is human LRR protein. The protein is used in the exemplification of the
CC invention.
XX
SQ Sequence 635 AA;

Query Match 56.8%; Score 42; DB 8; Length 635;
Best Local Similarity 63.6%; Pred. No. 61;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||
Db 432 GVSLSKLSLHN 442

RESULT 10
AAY94963
ID AAY94963 standard; protein; 695 AA.
XX
AC AAY94963;
XX
DT 16-JUN-2000 (first entry)
XX
DE Human secreted protein clone nf56_3 protein sequence SEQ ID NO:132.
XX
KW Human; secreted protein; immunestimulant; immunesuppressant; virucide;
KW antibacterial; antifungal; cytostatic; antiinflammatory; dermatological;

KW antidiabetic; antiasthmatic; antiarthritic; antirheumatic; protozoacide;
KW antithyroid; immune deficiency; severe combined immunedeficiency; SCID;
KW infection; HIV; hepatitis; malaria; autoimmune disorder; systemic lupus;
KW connective tissue disease; multiple sclerosis; erythematosis;
KW rheumatoid arthritis; autoimmune pulmonary inflammation; asthma;
KW Guillain-Barre syndrome; autoimmune thyroiditis; myasthenia gravis;
KW insulin dependent diabetes mellitus; graft-versus-host-disease;
KW autoimmune inflammatory eye disease; allergy.
XX
OS Homo sapiens.
XX
PN WO200009552-A1.
XX
PD 24-FEB-2000.
XX
PF 13-AUG-1999; 99WO-US018298.
XX
PR 14-AUG-1998; 98US-0096622P.
PR 17-AUG-1998; 98US-0096815P.
PR 04-SEP-1998; 98US-0099229P.
PR 23-OCT-1998; 98US-0105368P.
PR 08-JAN-1999; 99US-0115234P.
PR 12-FEB-1999; 99US-0119931P.
PR 18-FEB-1999; 99US-0120575P.
PR 30-APR-1999; 99US-0132020P.
PR 11-AUG-1999; 99US-0148424P.
XX
PA (GEMY) GENETICS INST INC.
XX
PI Jacobs K, Mccoy JM, Lavallie ER, Collins-Racie LA, Evans C;
PI Merberg D, Treacy M, Agostino MJ, Steininger RJ, Spaulding V;
PI Wong GG, Clark HF, Fechtel K;
XX
DR WPI; 2000-205979/18.
XX
PT New polynucleotides encoding secreted proteins, which may have e.g.
PT nutritional, chemokine, immune stimulating or suppressing, hematopoiesis
PT regulating, tissue growth, activin/inhibin antiinflammatory or tumor
PT inhibition activity.
XX
PS Claim 141; Page 592-594; 641pp; English.
XX
CC AAA16618 to AAA16697 encode the human secreted proteins given in AAY94898
CC to AAY94980, isolated from human adult brain, adult thyroid, adult
CC retina, foetal carcinoma, adult blood, adult neural, foetal kidney, adult
CC placenta, adult testis, whole embryo, adult cartilage, kidney, foetal
CC brain, adult thymus, foetal placenta, adult uterus, adult tumour, and
CC adult bladder, cDNA libraries. The polynucleotides and proteins are
CC predicted to have biological activities which would make them suitable
CC for treating, preventing or ameliorating medical conditions in humans and
CC animals. The polynucleotides can be used as markers for tissues in which
CC the protein is preferentially expressed, as molecular weight markers on
CC Southern gels, and as chromosome markers or tags to identify chromosomes
CC or to map gene positions. The proteins can be used in the treatment of
CC immune deficiencies and disorders, such as severe combined
CC immunedeficiency (SCID), as well as viral, bacterial, fungal and other
CC infections. These infections include human immunodeficiency virus (HIV),
CC hepatitis, herpesviruses, mycobacteria, Leismania spp., malaria and
CC candidiasis. The proteins can be used to treat autoimmune disorders such
CC as connective tissue disease, multiple sclerosis, systemic lupus
CC erythematosis, rheumatoid arthritis, autoimmune pulmonary inflammation,
CC Guillain-Barre syndrome, autoimmune thyroiditis, insulin dependent
CC diabetes mellitus, myasthenia gravis, graft-versus-host-disease and
CC autoimmune inflammatory eye disease. The proteins can also be used to
CC treat allergic conditions, such asthma. AAA16698 to AAA16774 represent
CC probes for the human secreted proteins from the present invention
XX
SQ Sequence 695 AA;

Query Match 56.8%; Score 42; DB 3; Length 695;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||
Db 492 GVSLSKLSLHN 502

RESULT 11
AAY08076
ID AAY08076 standard; protein; 696 AA.
XX
AC AAY08076;
XX
DT 11-SEP-2000 (first entry)
XX
DE Human PRO266 clone UNQ233 derived protein.
XX
KW Inflammatory cell infiltration; immune response; T cell proliferation;
KW anti-inflammatory; anti-autoimmune; anti-diabetic; spondyloarthropathy;
KW T cell-mediated disease; spondyloarthropathy; sclerosis; renal disease;
KW inflammatory myopathy; hemolytic anemia; thrombocytopenia; thyroiditis;
KW diabetes mellitus; demyelinating polyneuropathy; Guillain-Barre syndrome;
KW multiple sclerosis; polyneuropathy; hepatitis; cirrhosis; enteropathy;
KW sclerosing cholangitis; inflammatory bowel disease; Whipple's disease;
KW skin disease; dermatitis; psoriasis; asthma; allergic rhinitis; tumor;
KW food hypersensitivity; urticaria; eosinophilic pneumonia; transplant;
KW idiopathic pulmonary fibrosis; graft rejection; PRO245; human; PRO266;
KW UNQ233.
XX
OS Homo sapiens.
XX
PN WO9914241-A2.
XX
PD 25-MAR-1999.
XX
PF 17-SEP-1998; 98WO-US019437.
XX
PR 17-SEP-1997; 97US-0059119P.
PR 18-SEP-1997; 97US-0059263P.
PR 28-OCT-1997; 97US-0063550P.
PR 12-NOV-1997; 97US-0065186P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066770P.
PR 04-JUN-1998; 98US-0088026P.
XX
PA (GETH) GENENTECH INC.
XX
PI Fong S, Goddard A, Gurney AL, Tumas D, Wood WI;
XX
DR WPI; 1999-229499/19.
DR N-PSDB; AAX37722.
XX
PT Composition containing novel polypeptide PRO245, its agonist or
PT antagonist.
XX
PS Example 1; Fig 21; 177pp; English.
XX
CC This invention describes a novel composition containing (apart from a
CC carrier or excipient), a novel PRO245 polypeptide (I), its agonist or
CC antagonist, or their fragments, for modulating: (i) infiltration of
CC inflammatory cells into tissue; (ii) an immune response; or (iii) T cell
CC proliferation. The composition increases or decreases any of the effects
CC (i)-(iii). The products of the invention have anti-inflammatory, anti-
CC autoimmune and anti-diabetic activity. (I), and its (ant)agonists and
CC their fragments, are used to treat immune-related diseases, particularly
CC T cell-mediated diseases. The diseases treated include systemic lupus
CC erythematosus, rheumatoid arthritis, juvenile chronic arthritis,
CC spondyloarthropathies, systemic sclerosis (scleroderma), idiopathic
CC inflammatory myopathies (dermatomyositis, polymyositis), Sjogren's
CC syndrome, systemic vasculitis, sarcoidosis, autoimmune hemolytic anemia
CC (immune pancytopenia, paroxysmal nocturnal hemoglobinuria), autoimmune
CC thrombocytopenia (idiopathic thrombocytopenic purpura immune-mediated
CC thrombocytopenia), thyroiditis (Grave's disease, Hashimoto's thyroiditis,
CC juvenile lymphocytic thyroiditis, atrophic thyroiditis), diabetes
CC mellitus, immune-mediated renal disease (glomerulonephritis,

```
CC   tubulointerstitial nephritis), multiple sclerosis, idiopathic
CC   demyelinating polyneuropathy, Guillain-Barre syndrome, chronic
CC   inflammatory demyelinating polyneuropathy, infectious hepatitis
CC   (hepatitis A, B, C, D, E and other non-hepatotropic viruses), autoimmune
CC   chronic active hepatitis, primary biliary cirrhosis, granulomatous
CC   hepatitis, and sclerosing cholangitis, inflammatory bowel disease
CC   (ulcerative colitis: Crohn's disease), gluten-sensitive enteropathy, and
CC   Whipple's disease. Autoimmune or immune-mediated skin diseases including
CC   bullous skin diseases, erythema multiforme,contact dermatitis, psoriasis,
CC   asthma, allergic rhinitis, atopic dermatitis, food hypersensitivity,
CC   urticaria, eosinophilic pneumonia, idiopathic pulmonary fibrosis,
CC   hypersensitivity pneumonitis, and transplantation associated diseases
CC   (graft rejection, and graft-versus-host-disease). (I), its (ant)agonists
CC   or fragment can also be used as an adjuvant in treatment of tumors.
CC   Antibodies against (I) can also be used for diagnosing such diseases.
CC   This sequence represents a protein derived from human PRO266 clone UNQ233
CC   cDNA which is described in the method of the invention
XX
SQ   Sequence 696 AA;

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 12
AAY13359
ID   AAY13359 standard; protein; 696 AA.
XX
AC   AAY13359;
XX
DT   25-JUN-1999  (first entry)
XX
DE   Amino acid sequence of protein PRO266.
XX
KW   Secreted protein; transmembrane protein; human; enterocolitis;
KW   Zollinger-Ellison syndrome; gastrointestinal ulceration;
KW   congenital microvillus atrophy; skin disease; cell growth;
KW   abnormal keratinocyte differentiation; psoriasis; epithelial cancer;
KW   Parkinson's disease; Alzheimer's disease; ALS; neuropathy; fibromodulin;
KW   dermal scarring; Usher Syndrome; Atrophia areata; anti-thrombotic;
KW   wound healing; tissue repair.
XX
OS   Homo sapiens.
XX
PN   WO9914328-A2.
XX
PD   25-MAR-1999.
XX
PF   16-SEP-1998;   98WO-US019330.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
PR   24-OCT-1997;   97US-0063120P.
PR   24-OCT-1997;   97US-0063121P.
PR   24-OCT-1997;   97US-0063127P.
PR   24-OCT-1997;   97US-0063128P.
PR   27-OCT-1997;   97US-0063327P.
PR   27-OCT-1997;   97US-0063329P.
PR   28-OCT-1997;   97US-0063541P.
PR   28-OCT-1997;   97US-0063542P.
PR   28-OCT-1997;   97US-0063544P.
PR   28-OCT-1997;   97US-0063549P.
PR   28-OCT-1997;   97US-0063550P.
PR   28-OCT-1997;   97US-0063564P.
PR   29-OCT-1997;   97US-0063435P.
PR   29-OCT-1997;   97US-0063704P.
PR   29-OCT-1997;   97US-0063732P.
PR   29-OCT-1997;   97US-0063734P.
PR   29-OCT-1997;   97US-0063735P.
PR   29-OCT-1997;   97US-0063738P.
PR   29-OCT-1997;   97US-0064215P.
PR   31-OCT-1997;   97US-0063870P.
PR   31-OCT-1997;   97US-0064103P.
PR   03-NOV-1997;   97US-0064248P.
PR   07-NOV-1997;   97US-0064809P.
PR   12-NOV-1997;   97US-0065186P.
PR   17-NOV-1997;   97US-0065846P.
PR   18-NOV-1997;   97US-0065693P.
PR   21-NOV-1997;   97US-0066120P.
PR   21-NOV-1997;   97US-0066364P.
PR   24-NOV-1997;   97US-0066453P.
PR   24-NOV-1997;   97US-0066466P.
PR   24-NOV-1997;   97US-0066511P.
PR   24-NOV-1997;   97US-0066770P.
PR   24-NOV-1997;   97US-0066772P.
PR   25-NOV-1997;   97US-0066840P.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Wood WI,  Gurney AL,  Goddard A,  Pennica D,  Chen J,  Yuan J;
XX
DR   WPI; 1999-229533/19.
DR   N-PSDB; AAX52230.
XX
PT   New isolated human genes and polypeptides used in, e.g. treatment of
PT   gastrointestinal ulceration.
XX
PS   Claim 12; Fig 34; 320pp; English.
XX
CC   AAY13344-403 represent secreted and transmembrane human proteins. The
CC   cDNA sequences are obtained from cDNA libraries, prepared from fetal
CC   lung, fetal kidney, fetal brain, fetal liver and fetal retina. The
CC   encoded polypeptides have specific uses based on their homology to known
CC   polypeptides, e.g. PRO211 and PRO217 can be used for disorders associated
CC   with the preservation and maintenance of gastrointestinal mucosa and the
CC   repair of acute and chronic mucosal lesions (e.g. enterocolitis,
CC   Zollinger-Ellison syndrome, gastrointestinal ulceration and congenital
CC   microvillus atrophy), skin diseases associated with abnormal keratinocyte
CC   differentiation (e.g. psoriasis, epithelial cancers such as lung squamous
CC   cell carcinoma of the vulva and gliomas), potent effects on cell growth
CC   and development, diseases related to growth or survival of nerve cells
CC   including Parkinson's disease, Alzheimer's disease, ALS, neuropathies or
CC   cancer. PRO265 can be used as for fibromodulin, e.g. for reducing dermal
CC   scarring. PRO264 can be used as a target for anti-tumor drugs. PRO533 may
CC   be used in the treatment of Usher Syndrome or Atrophia areata; PRO269 can
CC   be used as an anti-thrombotic agent; PRO287 polypeptides and portions may
CC   have therapeutic applications in wound healing and tissue repair; PRO317
CC   can be used for treating problems of the kidney, uterus, endometrium,
CC   blood vessels, or related tissue, e.g. in the heart of genital tract
XX
SQ   Sequence 696 AA;

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
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              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 13
AAY70671
ID   AAY70671 standard; protein; 696 AA.
XX
AC   AAY70671;
XX
DT   18-JUL-2000  (first entry)
XX
DE   Human PRO266 protein.
XX
KW   PRO266; UNQ233; dermatological; immunosuppressive; antiinflammatory;
KW   immunostimulant; antiasthmatic; antirheumatic; antiarthritic; virucide;
KW   antiallergic; haemostatic; hepatotropic; antidiabetic; antianaemic;
KW   nephrotropic; neuroprotective; anticoagulant; immunological disorder;
KW   lung; pneumonia; skin; psoriasis; kidney; glomerulonephritis; arthritis;
KW   spondyloarthropathy; SLE; systemic lupus erythematosis; scleroderma;
KW   idiopathic inflammatory myopathy; anaemia; thrombocytopenia; diabetes;
KW   thyroiditis; Grave's disease; demyelinating disease; multiple sclerosis;
KW   Crohn's disease; hepatobiliary disease; hepatitis; asthma; human;
KW   graft-versus-host-disease.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
FT   Modified-site   17. .23
FT                   /note= "N-myristoylation site"
FT   Modified-site   18. .22
FT                   /note= "N-glycosylation site"
FT   Modified-site   30. .34
FT                   /note= "Casein Kinase II phosphorylation site"
FT   Modified-site   67. .73
FT                   /note= "N-myristoylation site"
FT   Modified-site   100. .106
FT                   /note= "N-myristoylation site"
FT   Modified-site   122. .126
FT                   /note= "cAMP and cGMP-dependent protein kinase
FT                   phosphorylation site"
FT   Modified-site   180. .184
FT                   /note= "Casein Kinase II phosphorylation site"
FT   Modified-site   222. .226
FT                   /note= "Casein Kinase II phosphorylation site"
FT   Modified-site   253. .257
FT                   /note= "N-glycosylation site"
FT   Modified-site   256. .260
FT                   /note= "Casein Kinase II phosphorylation site"
FT   Modified-site   302. .308
FT                   /note= "N-myristoylation site"
FT   Modified-site   328. .334
FT                   /note= "N-myristoylation site"
FT   Binding-site    337. .348
FT                   /note= "Prokaryotic membrane lipoprotein lipid attachment
FT                   site"
FT   Modified-site   343. .349
FT                   /note= "N-myristoylation site"
FT   Modified-site   354. .360
FT                   /note= "N-myristoylation site"
FT   Modified-site   363. .367
FT                   /note= "N-glycosylation site"
FT   Modified-site   366. .370
FT                   /note= "Casein Kinase II phosphorylation site"
FT   Modified-site   416. .420
FT                   /note= "N-glycosylation site"
FT   Modified-site   465. .471
FT                   /note= "N-myristoylation site"
FT   Modified-site   493. .499
FT                   /note= "N-myristoylation site"
FT   Modified-site   573. .577
FT                   /note= "Casein Kinase II phosphorylation site"
FT   Modified-site   595. .599
FT                   /note= "N-glycosylation site"
FT   Modified-site   598. .604
FT                   /note= "N-myristoylation site"
FT   Modified-site   603. .609
FT                   /note= "N-myristoylation site"
FT   Modified-site   608. .612
FT                   /note= "Casein Kinase II phosphorylation site"
FT   Modified-site   646. .650
FT                   /note= "cAMP and cGMP-dependent protein kinase
FT                   phosphorylation site"
FT   Modified-site   655. .659
FT                   /note= "N-glycosylation site"
FT   Modified-site   657. .661
FT                   /note= "Casein Kinase II phosphorylation site"
FT   Modified-site   666. .670
FT                   /note= "Casein Kinase II phosphorylation site"
FT   Modified-site   693. .697
FT                   /note= "Casein Kinase II phosphorylation site"
XX
PN   WO200015797-A2.
XX
PD   23-MAR-2000.
XX
PF   15-SEP-1999;   99WO-US021547.
XX
PR   17-SEP-1998;   98US-0100858P.
PR   17-SEP-1998;   98WO-US019437.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Fong S,  Goddard A,  Gurney AL,  Tumas D,  Wood WI;
XX
DR   WPI; 2000-271435/23.
DR   N-PSDB; AAZ52205.
XX
PT   Composition for treatment and diagnosis of immune related diseases e.g.
PT   Grave's disease comprises a PRO245, PRO217, PRO301, PRO266, PRO335,
PT   PRO331 or PRO326 polypeptide or its agonists or antagonists (preferably
PT   antibodies).
XX
PS   Example 1; Fig 10; 201pp; English.
XX
CC   The present sequence is the human protein PRO266, encoded by UNQ233 cDNA,
CC   designated as clone DNA37150. It is isolated from human foetal brain
CC   tissue. PRO266 has significant homology to a SLIT protein, indicating
CC   that it could be a leucine rich repeat protein. It enhances or suppresses
CC   the infiltration of inflammatory cells into tissues, proliferation of T-
CC   lymphocytes and modulates the immune response. This sequence is useful
CC   for treatment of immune related disorders, like SLE, rheumatoid/juvenile
CC   arthritis, spondyloarthropathy, systemic sclerosis (scleroderma),
CC   idiopathic inflammatory myopathies such as dermatomyositis, Sjogren's
CC   syndrome, systemic vasculitis, sarcoidosis, autoimmune haemolytic
CC   anaemia, thrombocytopenia, thyroiditis e.g. Grave's disease, diabetes
CC   mellitus, immune-mediated renal disease e.g. glomerulonephritis,
CC   demyelinating diseases such as multiple sclerosis and Guillain-Barre
CC   syndrome, hepatobiliary diseases like hepatitis and primary biliary
CC   cirrhosis, inflammatory and fibrotic lung diseases such as inflammatory
CC   bowel disease (e.g. Crohn's disease), autoimmune or immune-mediated skin
CC   diseases such as psoriasis, allergies like asthma, immunological diseases
CC   of the lungs such as eosinophilic pneumonia and transplantation
CC   associated diseases such as graft-versus-host-disease
XX
SQ   Sequence 696 AA;

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

```
RESULT 14
ADC78411
ID   ADC78411 standard; protein; 696 AA.
XX
AC   ADC78411;
XX
DT   01-JAN-2004  (first entry)
XX
DE   Human PRO266 protein.
XX
KW   antiinflamatory; antiulcer; cytostatic; antipsoriatic; antiparkinsonian;
KW   nootropic; neuroprotective; vasotropic; chemotaxic; angiogenic;
KW   neurotrophic; osteopathic; antiasthmatic; antiarthritic; antirheumatic;
KW   antiarteriosclerotic; cardiant; antidiabetic; cerebroprotective;
KW   thrombolytic; immunomodulator; enterocolitis; Zollinger-Ellison syndrome;
KW   gastrointestinal ulceration; psoriasis; cancer; Parkinson's disease;
KW   Alzheimer's; ALS; neuropathy; dermal scarring; wound healing;
KW   nerve repair; thrombosis; bone; cartilage formation; angiogenesis;
KW   asthma; rheumatoid arthritis; multiple sclerosis; inflammatory disorder;
KW   atherosclerosis; cardiac injury; infertility; premature aging; AIDS;
KW   diabetes; stroke; gene therapy; transgenic; PRO; human.
XX
OS   Homo sapiens.
XX
PN   WO200015796-A2.
XX
PD   23-MAR-2000.
XX
PF   15-SEP-1999;   99WO-US021090.
XX
PR   16-SEP-1998;   98WO-US019330.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Chen J,  Goddard A,  Gurney AL,  Hillan K,  Pennica D,  Wood WI;
PI   Yuan J;
XX
DR   WPI; 2000-271434/23.
DR   N-PSDB; ADC78410.
XX
PT   Novel nucleic acids encoding secreted and transmembrane polypeptides with
PT   homology, e.g. to growth and cancer-associated antigens.
XX
PS   Claim 12; SEQ ID NO 91; 355pp; English.
XX
CC   The invention relates to a novel nucleic acid encoding a PRO polypeptide.
CC   The polypeptides and polynucleotides of the invention may be useful as
CC   research tools and as therapeutics for treating enterocolitis, Zollinger-
CC   Ellison syndrome, gastrointestinal ulceration, psoriasis, cancer,
CC   Parkinson's disease, Alzheimer's disease, ALS, neuropathies, dermal
CC   scarring and wound healing, nerve repair, thrombosis, bone and/or
CC   cartilage formation, angiogenesis, asthma, rheumatoid arthritis, multiple
CC   sclerosis, inflammatory disorders, atherosclerosis, cardiac injury,
CC   infertility, premature aging, AIDS, diabetes complications and stroke.
CC   The molecules may also be utilised during gene therapy procedures and
CC   transgenic animal production. The current sequence is that of the human
CC   PRO protein of the invention.
XX
SQ   Sequence 696 AA;

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 15
AAB80227
ID   AAB80227 standard; protein; 696 AA.
XX
AC   AAB80227;
XX
DT   24-APR-2001  (first entry)
XX
DE   Human PRO266 protein.
XX
KW   Human; PRO; dermatological; antipsoriatic; cytostatic; antiinflammatory;
KW   antiparkinsonian nootropic; neuroprotective; vulnerary; cardiant;
KW   antiangiogenic; vasotropic; antiasthmatic; antirheumatic; cancer;
KW   antiarthritic; antiinfertility; antidiabetic; antiviral; diabetes;
KW   ophthalmological; gene therapy; skin disease; gastrointestinal disorder;
KW   ischaemia; inflammation.
XX
OS   Homo sapiens.
XX
PN   WO200104311-A1.
XX
PD   18-JAN-2001.
XX
PF   22-FEB-2000; 2000WO-US004414.
XX
PR   07-JUL-1999;   99US-0143048P.
PR   26-JUL-1999;   99US-0145698P.
PR   28-JUL-1999;   99US-0146222P.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   02-DEC-1999;   99WO-US028564.
PR   02-DEC-1999;   99WO-US028565.
PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   05-JAN-2000; 2000WO-US000219.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Ashkenazi AJ,  Botstein D,  Desnoyers L,  Eaton DL,  Ferrara N;
PI   Filvaroff E,  Fong S,  Gao W,  Gerber H,  Gerritsen ME,  Goddard A;
PI   Godowski PJ,  Grimaldi CJ,  Gurney AL,  Hillan KJ,  Kljavin IJ;
PI   Mather JP,  Pan J,  Paoni NF,  Roy MA,  Stewart TA,  Tumas D;
PI   Williams PM,  Wood WI;
XX
DR   WPI; 2001-081051/09.
DR   N-PSDB; AAF72388.
XX
PT   Sixty one nucleic acids encoding PRO polypeptides which are useful in the
PT   treatment of skin diseases (e.g. psoriasis), cancers (e.g. lung squamous
PT   cell carcinoma) and neurodegenerative diseases (e.g. Alzheimer's
PT   disease).
XX
PS   Claim 1; Fig 34; 393pp; English.
XX
CC   The present sequence is one of sixty one novel secreted and transmembrane
CC   PRO polypeptides. The PRO polypeptides are useful for treating skin
CC   diseases (e.g. psoriasis), cancers (e.g. lung squamous cell carcinoma),
CC   gastrointestinal disorders (e.g. enterocolitis), neurodegenerative
CC   diseases (e.g. Alzheimer's disease, Parkinson's disease), wound repair,
CC   cardiovascular disorders (e.g. endometrial bleeding angiogenesis,
CC   ischaemias such as coronary ischaemia, atherosclerosis), inflammatory
CC   disorders (e.g. asthma, rheumatoid arthritis, multiple sclerosis),
CC   infertility, AIDS and diabetes and retinal disorders such as retinitis
CC   pigmentosum. The PRO nucleic acids have applications in molecular
CC   biology, including use as hybridization probes, and in chromosome and
CC   gene mapping
XX
SQ   Sequence 696 AA;

  Query Match             56.8%;  Score 42;  DB 4;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
```

```
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 16
AAU00824
ID   AAU00824 standard; protein; 696 AA.
XX
AC   AAU00824;
XX
DT   04-JUL-2001  (first entry)
XX
DE   Human immune response protein PRO266 (UNQ233).
XX
KW   Human; PRO266; UNQ233; immune response; osteoarthritis;
KW   systemic lupus erythematosus; rheumatoid arthritis; systemic sclerosis;
KW   juvenile chronic arthritis; spondyloarthropathy; Sjogren's syndrome;
KW   idiopathic inflammatory; myopathy; polymyositis; systemic vasculitis;
KW   sarcoidosis; autoimmune haemolytic anaemia; immune pancytopaenia;
KW   autoimmune thrombocytopaenia; idiopathic thrombocytopaenic purpura;
KW   thyroiditis; Grave's disease; Hashimoto's thyroiditis; diabetes mellitus;
KW   glomerulonephritis; demyelinating disease; multiple sclerosis;
KW   Guillain-Barre syndrome; hepatobiliary disease;
KW   chronic inflammatory demyelinating polyneuropathy; infectious hepatitis;
KW   auto immune chronic active hepatitis; primary biliary cirrhosis;
KW   granulomatous hepatitis; sclerosing cholangitis; ulcerative colitis;
KW   inflammatory bowel disease; Crohn's disease; Whipple's disease;
KW   erythaemia multiforme; psoriasis; asthma; allergic rhinitis; urticaria;
KW   food hypersensitivity; eosinophilic pneumonia; graft rejection;
KW   idiopathic pulmonary fibrosis; graft-versus-host-disease; immunogen;
KW   antibody.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
FT   Peptide         1. .15
FT                   /label= Signal_peptide
FT   Protein         16. .696
FT                   /label= Mature_PRO266
FT   Modified-site   17. .23
FT                   /note= "Glycine at 17 is N-myristoylated"
FT   Modified-site   18. .22
FT                   /note= "Asn is N-glycosylated"
FT   Modified-site   67. .73
FT                   /note= "Glycine is N-myristoylated"
FT   Modified-site   100. .106
FT                   /note= "Glycine at 100 is N-myristoylated"
FT   Region          122. .126
FT                   /label= Phosphorylation_site
FT                   /note= "cAMP/cGMP dependent protein kinase
FT                   phosphorylation site"
FT   Modified-site   253. .257
FT                   /note= "Asn is N-glycosylated"
FT   Modified-site   302. .308
FT                   /note= "Glycine at 302 is N-myristoylated"
FT   Modified-site   328. .334
FT                   /note= "Glycine is N-myristoylated"
FT   Region          337. .348
FT                   /label= Lipid_attachment_site
FT                   /note= "Prokaryotic membrane lipoprotein attachment site"
FT   Modified-site   343. .349
FT                   /note= "Glycine at 343 is N-myristoylated"
FT   Modified-site   354. .360
FT                   /note= "Glycine is N-myristoylated"
FT   Modified-site   363. .367
FT                   /note= "Asn is N-glycosylated"
FT   Modified-site   416. .420
FT                   /note= "Asn is N-glycosylated"
FT   Modified-site   465. .471
FT                   /note= "Glycine is N-myristoylated"
FT   Modified-site   493. .499
FT                   /note= "Glycine is N-myristoylated"
FT   Modified-site   595. .599
FT                   /note= "Asn is N-glycosylated"
FT   Modified-site   598. .604
FT                   /note= "Glycine at 598 is N-myristoylated"
FT   Modified-site   603. .609
FT                   /note= "Glycine is N-myristoylated"
FT   Domain          619. .639
FT                   /label= Transmembrane_domain
FT   Modified-site   655. .659
FT                   /note= "Asn is N-glycosylated"
XX
PN   WO200119991-A1.
XX
PD   22-MAR-2001.
XX
PF   20-MAR-2000; 2000WO-US007377.
XX
PR   15-SEP-1999;   99WO-US021547.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Fong S,  Goddard A,  Gurney AL,  Hillan KJ,  Tumas D,  Wood WI;
XX
DR   WPI; 2001-226823/23.
DR   N-PSDB; AAS00160.
XX
PT   Composition for diagnosing and treating immune related diseases, e.g.
PT   rheumatoid arthritis and diabetes mellitus, comprises a PRO polypeptide,
PT   agonist, antagonist or fragment.
XX
PS   Claim 31; Fig 8; 138pp; English.
XX
CC   The sequence represents Human PRO266 (UNQ233), a protein involved in the
CC   immune response. PRO polypeptides, and (ant)agonists to them, are used in
CC   compositions for modulating infiltration of inflammatory cells into a
CC   tissue, modulating an immune response and modulating proliferation of T-
CC   lymphocytes in response to an antigen. Immune related diseases can be
CC   treated with the compositions, such as, systemic lupus erythematosus,
CC   rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis,
CC   spondyloarthopathies, systemic sclerosis, idiopathic inflammatory
CC   myopathies (e.g. polymyositis), Sjogren's syndrome, systemic vasculitis,
CC   sarcoidosis, autoimmune haemolytic anaemia (e.g. immune pancytopaenia),
CC   autoimmune thrombocytopaenia (e.g. idiopathic thrombocytopaenic purpura),
CC   thyroiditis (e.g. Grave's disease, Hashimoto's thyroiditis), diabetes
CC   mellitus, immune-mediated renal disease (e.g. glomerulonephritis),
CC   demyelinating diseases of the central and peripheral nervous systems e.g.
CC   multiple sclerosis or Guillain-Barre syndrome, and chronic inflammatory
CC   demyelinating polyneuropathy, hepatobiliary diseases such as infectious
CC   hepatitis (hepatitis A, B, C, D, E and other non-hepatotropic viruses),
CC   auto immune chronic active hepatitis, primary biliary cirrhosis,
CC   granulomatous hepatitis, and sclerosing cholangitis, inflammatory bowel
CC   disease (ulcerative colitis, Crohn's disease and Whipple's disease),
CC   autoimmune or immune-mediated skin diseases (e.g. erythaemia multiforme
CC   and psoriasis), asthma, allergic rhinitis, urticaria, food
CC   hypersensitivity, immunologic diseases of the lung such as eosinophilic
CC   pneumonias, idiopathic pulmonary fibrosis, transplantation associated
CC   diseases including graft-versus-host-disease and graft rejection. PRO
CC   polypeptides can be used to diagnose immune related diseases, to identify
CC   inhibitors, and to stimulate the proliferation of T lymphocytes. Anti-PRO
CC   antibodies can be used to detect PRO and in diagnosis. PRO polypeptides,
CC   antibodies and (ant)agonists can be used in rational drug design
XX
SQ   Sequence 696 AA;

  Query Match             56.8%;  Score 42;  DB 4;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 17
AAU12348
ID AAU12348 standard; protein; 696 AA.
XX
AC AAU12348;
XX
DT 24-OCT-2001 (first entry)
XX
DE Human PRO266 polypeptide sequence.
XX
KW Human secretory and transmembrane; PRO; mammalian; cancer; lung; breast;
KW prostate; cervical; tumour necrosis factor-alpha; TNF-alpha; cartilage;
KW ear; proliferation; glucose; free fatty acid; skeletal muscle; adipocyte;
KW A-peptide; factor VIIA; gene therapy.
XX
OS Homo sapiens.
XX
PN WO200140466-A2.
XX
PD 07-JUN-2001.
XX
PF 01-DEC-2000; 2000WO-US032678.
XX
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 02-DEC-1999; 99WO-US028551.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 09-DEC-1999; 99US-0170262P.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 30-DEC-1999; 99WO-US031243.
PR 30-DEC-1999; 99WO-US031274.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000277.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 18-FEB-2000; 2000WO-US004342.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 01-MAR-2000; 2000WO-US005601.
PR 02-MAR-2000; 2000WO-US005841.
PR 03-MAR-2000; 2000US-0187202P.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
PR 21-MAR-2000; 2000WO-US007532.
PR 30-MAR-2000; 2000WO-US008439.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 05-JUN-2000; 2000US-0209832P.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 10-NOV-2000; 2000WO-US030873.
XX
PA (GETH) GENENTECH INC.
XX
PI Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX
DR WPI; 2001-408281/43.
DR N-PSDB; AAS21420.
XX
PT Isolated , secretory and transmembrane PRO polypeptide used to detect
PT other PRO polypeptides, link bioactive molecules to cells expressing PRO
PT polypeptides, and detect the presence of mammalian tumors e.g. lung,
PT breast, prostate, cervical.
XX
PS Claim 12; Fig 354; 813pp; English.
XX
CC AAU12172-AAU12446 represent novel human secretory and transmembrane PRO
CC polypeptides. The PRO polypeptides are useful to detect other PRO
CC polypeptides, to link bioactive molecules to cells expressing PRO
CC polypeptides, to modulate biological activities of cells expressing PRO
CC polypeptides, and to detect the presence of mammalian lung, colon,
CC breast, prostate, rectal, cervical or liver tumours by comparing PRO
CC polypeptide expression in a cell sample to that in a control sample. Some
CC of the 275 sequences are also useful to stimulate the release of tumour
CC necrosis factor-alpha (TNF-alpha) from human blood, the proliferation or
CC differentiation of chondrocytes, the proliferation or gene expression in
CC pericyte cells, the release of proteoglycans from cartilage, the
CC proliferation of inner ear utricular supporting cells or of T-
CC lymphocytes, the release of a cytokine from peripheral blood monocytes
CC (PBMCs), or the proliferation of endothelial cells. Some of the PRO
CC polypeptides may modulate glucose or free fatty acid uptake by skeletal
CC muscle cells or by adipocytes; or inhibit binding of A-peptide to factor
CC VIIA. The PRO polypeptides can be used in assays to identify molecules
CC involved in binding interactions. The polynucleotides encoding PRO
CC polypeptides can be used to generate probes, antisense RNA/DNA,
CC transgenic or knock out animals and can be used in gene therapy
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 4; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 18
AAB50905
ID AAB50905 standard; protein; 696 AA.
XX
AC AAB50905;
XX
DT 21-MAR-2001 (first entry)
XX
DE Human PRO266 protein.
XX
KW Human; PRO; antiinflammatory; dermatological; antiarthritic;
KW antirheumatic; cardiant; antianaemic; immunosuppressive; antithyroid;
KW antidiabetic; nootropic; neuroprotective; hepatotropic; virucide;
KW antiallergic; antiasthmatic; immune related disorder;
KW hepatobiliary disease; autoimmune disease; allergy.
XX
OS Homo sapiens.
XX
PN WO200073452-A2.
XX
PD 07-DEC-2000.
XX
PF 02-JUN-2000; 2000WO-US015264.
XX
PR 02-JUN-1999; 99WO-US012252.
PR 20-JUL-1999; 99US-0144732P.
PR 20-JUL-1999; 99US-0144758P.
PR 28-JUL-1999; 99US-0146222P.
PR 01-SEP-1999; 99WO-US020111.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 29-OCT-1999; 99US-0162506P.
PR 30-NOV-1999; 99WO-US028313.

```
PR   01-DEC-1999;   99WO-US028634.
PR   02-DEC-1999;   99WO-US028551.
PR   02-DEC-1999;   99WO-US028565.
PR   09-DEC-1999;   99US-0170262P.
PR   20-DEC-1999;   99WO-US030911.
PR   05-JAN-2000; 2000WO-US000219.
PR   06-JAN-2000; 2000WO-US000376.
PR   11-FEB-2000; 2000WO-US003565.
PR   18-FEB-2000; 2000WO-US004341.
PR   18-FEB-2000; 2000WO-US004342.
PR   22-FEB-2000; 2000WO-US004414.
PR   24-FEB-2000; 2000WO-US004914.
PR   24-FEB-2000; 2000WO-US005004.
PR   01-MAR-2000; 2000WO-US005601.
PR   02-MAR-2000; 2000WO-US005841.
PR   03-MAR-2000; 2000US-0187202P.
PR   15-MAR-2000; 2000WO-US006884.
PR   20-MAR-2000; 2000WO-US007377.
PR   21-MAR-2000; 2000WO-US007532.
PR   30-MAR-2000; 2000WO-US008439.
PR   17-MAY-2000; 2000WO-US013705.
PR   22-MAY-2000; 2000WO-US014042.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Ashkenazi AJ,  Baker KP,  Chan B,  Goddard A,  Godowski PJ;
PI   Gurney AL,  Hebert C,  Henzel W,  Kabakoff RC,  Shelton DL,  Tumas D;
PI   Watanabe CK,  Wood WI;
XX
DR   WPI; 2001-025253/03.
DR   N-PSDB; AAC91464.
XX
PT   Thirty three nucleic acids encoding PRO polypeptides which are useful in
PT   the diagnosis and treatment of immune related disorders, e.g. systemic
PT   lupus erythematosis, rheumatoid arthritis, osteoarthritis, thyroiditis
PT   and diabetes mellitus.
XX
PS   Claim 58; Fig 8; 218pp; English.
XX
CC   The present sequence is one of thirty three novel PRO polypeptides. The
CC   PRO polypeptides, anti-PRO antibodies, agonists and antagonists are
CC   useful for treating and diagnosing immune related disorders such as
CC   systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis,
CC   juvenile chronic arthritis, spondyloarthropathies, systemic sclerosis,
CC   idiopathic inflammatory myopathies, Sjogren's syndrome, systemic
CC   vasculitis, sarcoidosis, autoimmune haemolytic anaemia, autoimmune
CC   thrombocytopaenia, thyroiditis, diabetes mellitus, immune-mediated renal
CC   disease, demyelinating diseases of the central and peripheral nervous
CC   systems (such as multiple sclerosis, idiopathic demyelinating
CC   polyneuropathy or Guillain-Barre syndrome, and chronic inflammatory
CC   demyelinating polyneuropathy), hepatobiliary diseases (such as
CC   infectious, autoimmune chronic active hepatitis, primary biliary
CC   cirrhosis, granulomatous hepatitis and sclerosing cholangitis),
CC   inflammatory bowel disease, gluten-sensitive enteropathy and Whipple's
CC   disease, autoimmune or immune-mediated skin diseases (such as bullous
CC   skin diseases, erythema multiforme, contact dermatitis, psoriasis),
CC   allergic diseases such as asthma, allergic rhinitis, atopic dermatitis,
CC   food hypersensitivity and urticaria), immunological diseases of the lung
CC   (such as eosinophilic pneumonias, idiopathic pulmonary fibrosis and
CC   hypersensitivity pneumonitis), transplantation associated diseases
CC   including graft rejection and graft-versus-host diseases
XX
SQ   Sequence 696 AA;

  Query Match             56.8%;  Score 42;  DB 4;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

```
RESULT 19
ABU71605
ID   ABU71605 standard; protein; 696 AA.
XX
AC   ABU71605;
XX
DT   16-JUN-2003  (first entry)
XX
DE   Human PRO polypeptide #16.
XX
KW   Human; PRO; secreted polypeptide; transmembrane polypeptide;
KW   pathological disorder; cardiac insufficiency disorder; protein secretion;
KW   pancreas; diabetes; gastrointestinal mucosa; mucosal lesion; psoriasis;
KW   skin disease; keratinocyte differentiation; epithelial cancer; tumour;
KW   lung squamous cell carcinoma; epidermoid carcinoma; vulva; glioma;
KW   cytostatic; cardiant; endocrine; antidiabetic; gastrointestinal;
KW   antiulcer; dermatological; vulnerary.
XX
OS   Homo sapiens.
XX
PN   US2002146709-A1.
XX
PD   10-OCT-2002.
XX
PF   18-JUL-2001; 2001US-00909088.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
PR   24-OCT-1997;   97US-0063120P.
PR   24-OCT-1997;   97US-0063121P.
PR   24-OCT-1997;   97US-0063127P.
PR   24-OCT-1997;   97US-0063128P.
PR   27-OCT-1997;   97US-0063327P.
PR   27-OCT-1997;   97US-0063329P.
PR   28-OCT-1997;   97US-0063541P.
PR   28-OCT-1997;   97US-0063542P.
PR   28-OCT-1997;   97US-0063544P.
PR   28-OCT-1997;   97US-0063549P.
PR   28-OCT-1997;   97US-0063550P.
PR   28-OCT-1997;   97US-0063564P.
PR   29-OCT-1997;   97US-0063435P.
PR   29-OCT-1997;   97US-0063704P.
PR   29-OCT-1997;   97US-0063732P.
PR   29-OCT-1997;   97US-0063734P.
PR   29-OCT-1997;   97US-0063735P.
PR   29-OCT-1997;   97US-0063738P.
PR   29-OCT-1997;   97US-0064215P.
PR   31-OCT-1997;   97US-0063870P.
PR   31-OCT-1997;   97US-0064103P.
PR   03-NOV-1997;   97US-0064248P.
PR   07-NOV-1997;   97US-0064809P.
PR   12-NOV-1997;   97US-0065186P.
PR   17-NOV-1997;   97US-0065846P.
PR   18-NOV-1997;   97US-0065693P.
PR   21-NOV-1997;   97US-0066120P.
PR   21-NOV-1997;   97US-0066364P.
PR   24-NOV-1997;   97US-0066453P.
PR   24-NOV-1997;   97US-0066466P.
PR   24-NOV-1997;   97US-0066511P.
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PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-328338/31.
DR N-PSDB; ACA58970.
XX
PT Isolated nucleic acid useful for e.g., treating pathological disorders
PT encodes a secreted or transmembrane protein.
XX
PS Claim 12; Fig 34; 473pp; English.
XX
CC The invention relates to human PRO polypeptides (secreted or
CC transmembrane polypeptides) and the polynucleotides encoding them. The
CC PRO polypeptides and polynucleotides can be used in treating pathological
CC disorders and tumours, in therapeutic treatment of cardiac insufficiency
CC disorders and in therapeutic treatment of disorders involving protein
CC secretion by the pancreas, including diabetes. They can also be used in
CC treating disorders associated with the preservation and maintenance of
CC gastrointestinal mucosa and the repair of acute and chronic mucosal
CC lesions, and skin diseases associated with abnormal keratinocyte
CC differentiation (e.g., psoriases, epithelial cancers such as lung
CC squamous cell carcinoma, epidermoid carcinoma of the vulva and gliomas).
CC The sequences can be used as molecular markers for protein
CC electrophoresis purposes and can be utilised in protein-protein binding
CC assays, biochemical screening assays, immunoassays and cell-based assays.
CC This sequence represents a human PRO polypeptide of the invention
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||

Db 493 GVSLSKLSLHN 503

RESULT 20
ABO17792
ID ABO17792 standard; protein; 696 AA.
XX
AC ABO17792;
XX
DT 26-AUG-2003 (first entry)
XX
DE Novel human secreted and transmembrane protein PRO266.
XX
KW Human; secreted and transmembrane protein; PRO; antiinflammatory;
KW antiarteriosclerotic; cardiant; anti-infertility; anti-HIV; cytostatic;
KW antidiabetic; gene therapy; tumour necrosis factor (TNF)-alpha release;
KW TNF-alpha release; cell proliferation; cell differentiation;
KW gene expression modulator; proteoglycan release; cytokine release;
KW tumour; inflammatory disease; organ failure; atherosclerosis;
KW cardiac injury; infertility; birth defect; premature aging; AIDS;
KW acquired immunodeficiency syndrome; cancer; diabetic complication;
KW chromosome mapping; gene mapping; pharmaceutical; diagnostic; biosensor;
KW bioreactor; tissue typing.
XX
OS Homo sapiens.
XX
PN US2003032156-A1.
XX
PD 13-FEB-2003.
XX
PF 06-MAY-2002; 2002US-00140474.
XX
PR 31-MAR-1997; 97WO-US005230.
PR 12-JUN-1998; 98WO-US012456.
PR 14-JUL-1998; 98WO-US014552.
PR 28-AUG-1998; 98WO-US017888.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019093.
PR 14-SEP-1998; 98WO-US019094.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 07-OCT-1998; 98WO-US021141.
PR 29-OCT-1998; 98WO-US022991.
PR 29-OCT-1998; 98WO-US022992.
PR 20-NOV-1998; 98WO-US024855.
PR 01-DEC-1998; 98WO-US025108.
PR 05-JAN-1999; 99WO-US000106.
PR 08-MAR-1999; 99WO-US005028.
PR 10-MAR-1999; 99WO-US005190.
PR 20-APR-1999; 99WO-US008615.
PR 14-MAY-1999; 99WO-US010733.
PR 02-JUN-1999; 99WO-US012252.
PR 01-SEP-1999; 99WO-US020111.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 30-NOV-1999; 99WO-US028409.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 02-DEC-1999; 99WO-US028551.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 22-DEC-1999; 99WO-US030720.
PR 30-DEC-1999; 99WO-US031243.
PR 30-DEC-1999; 99WO-US031274.

PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000277.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 18-FEB-2000; 2000WO-US004342.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 01-MAR-2000; 2000WO-US005601.
PR 02-MAR-2000; 2000WO-US005746.
PR 02-MAR-2000; 2000WO-US005841.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
PR 21-MAR-2000; 2000WO-US007532.
PR 30-MAR-2000; 2000WO-US008439.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 10-NOV-2000; 2000WO-US030873.
PR 01-DEC-2000; 2000WO-US032678.
PR 20-DEC-2000; 2000US-00747259.
PR 20-DEC-2000; 2000WO-US034956.
PR 28-FEB-2001; 2001US-00796498.
PR 28-FEB-2001; 2001WO-US006520.
PR 01-MAR-2001; 2001WO-US006666.
PR 09-MAR-2001; 2001US-00802706.
PR 14-MAR-2001; 2001US-00808689.
PR 22-MAR-2001; 2001US-00816744.
PR 05-APR-2001; 2001US-00828366.
PR 10-MAY-2001; 2001US-00854208.
PR 10-MAY-2001; 2001US-00854280.
PR 18-MAY-2001; 2001US-00860216.
PR 25-MAY-2001; 2001US-00866028.
PR 25-MAY-2001; 2001US-00866034.
PR 25-MAY-2001; 2001WO-US017092.
PR 01-JUN-2001; 2001US-00872035.
PR 01-JUN-2001; 2001WO-US017800.
PR 05-JUN-2001; 2001US-00874503.
PR 14-JUN-2001; 2001US-00882636.
PR 19-JUN-2001; 2001US-00886342.
PR 20-JUN-2001; 2001WO-US019692.
PR 21-JUN-2001; 2001US-00887879.
PR 22-JUN-2001; 2001WO-US020116.
PR 29-JUN-2001; 2001WO-US021066.
PR 09-JUL-2001; 2001WO-US021735.
PR 18-JUL-2001; 2001US-00908827.
PR 06-AUG-2001; 2001US-00924419.
PR 09-AUG-2001; 2001US-00927796.
PR 16-AUG-2001; 2001US-00931836.
PR 19-DEC-2001; 2001US-00028072.
XX
PA (GETH) GENENTECH INC.
XX
PI Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX
DR WPI; 2003-341980/32.
DR N-PSDB; ACD24029.
XX
PT New secreted and transmembrane PRO nucleic acids, for treating
PT inflammation, organ failure, atherosclerosis, cardiac injury,
PT infertility, birth defects, premature aging, axquired immunodeficiency
PT syndrome (AIDS), or cancer.
XX
PS Claim 12; Fig 354; 660pp; English.
XX
CC The invention describes an isolated nucleic acid (I) comprising, or which
CC has 80 % sequence identity to, or the full-length coding sequence of, one
CC of 275 nucleotide sequences, and which encodes a corresponding
CC polypeptide selected from 275 amino acid sequences, where all sequences
CC are given in the specification. The polypeptide encoded by (I) is used to
CC detect PRO polypeptides, link a bioactive molecule to a cell expressing a
CC PRO polypeptide, modulate a biological activity of a cell, stimulate the
CC release of tumour necrosis factor (TNF)-alpha from human blood, modulate
CC the uptake of glucose or free fatty acid by cells, stimulate or inhibit
CC the proliferation or differentiation of cells or gene expression,
CC stimulate the release of proteoglycans, stimulate the release of cytokine
CC from peripheral blood mononuclear cells, inhibit the binding of A-peptide
CC to factor VIIA, or detect the presence of tumour in a mammal. The nucleic
CC acid and polypeptide encoded by it, are useful for treating inflammatory
CC diseases, organ failure, atherosclerosis, cardiac injury, infertility,
CC birth defects, premature aging, acquired immunodeficiency syndrome
CC (AIDS), cancer, or diabetic complications. The nucleic acid is useful as
CC hybridisation probes, in chromosome and gene mapping, and in generating
CC antisense RNA or DNA. The polypeptides are useful as pharmaceuticals,
CC diagnostics, biosensors or bioreactors. Both are useful in tissue typing.
CC This is the amino acid sequence of a novel human secreted and
CC transmembrane PRO polypeptide
XX
SQ Sequence 696 AA;

```
  Query Match             56.8%;  Score 42;  DB 6;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 21
ABU71460
ID ABU71460 standard; protein; 696 AA.
XX
AC ABU71460;
XX
DT 10-JUN-2003 (first entry)
XX
DE Human PRO polypeptide #16.
XX
KW Human; secreted and transmembrane protein; PRO polypeptide; cancer;
KW Alzheimer's disease; ischaemia; cytostatic; nootropic; vasotropic;
KW neuroprotective.
XX
OS Homo sapiens.
XX
PN US2002192659-A1.
XX
PD 19-DEC-2002.
XX
PF 10-JUL-2001; 2001US-00902853.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.

PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-361832/34.
DR N-PSDB; ACA58367.
XX
PT New isolated nucleic acid encoding a PRO polypeptide, e.g. PRO245 or
PT PRO1868, useful in molecular biology, chromosome and gene mapping, in
PT generating antisense RNA and DNA, and in gene therapy.
XX
PS Claim 12; Fig 34; 474pp; English.
XX
CC The present invention relates to the isolation of novel human secreted
CC and transmembrane proteins (PRO polypeptides), and the polynucleotide
CC sequences encoding them. The polynucleotide sequences are useful in
CC molecular biology, as hybridisation probes, in chromosome and gene
CC mapping, in generating antisense RNA and DNA, and in gene therapy. The
CC polynucleotide sequences may also be used in preparing PRO polypeptides
CC by recombinant techniques, and in generating either transgenic animals or
CC knock-out animals which, in turn, are useful in the development and
CC screening of therapeutically useful reagents. The PRO polypeptides or
CC their antibodies are useful in preparing a medicament for treating a
CC condition responsive to the polypeptide or antibody, such as cancer,
CC Alzheimer's disease or ischaemia, and in various diagnostic assays.
CC ABU71445-ABU71505 represent human PRO polypeptides of the invention
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 22
ABU81046
ID ABU81046 standard; protein; 696 AA.
XX
AC ABU81046;
XX
DT 23-JUN-2003 (first entry)
XX
DE Human PRO polypeptide #177.
XX
KW Human; PRO polypeptide; secreted and transmembrane protein;
KW anti-PRO antibody; diagnostic assay; gene expression; diabetes;
KW bone disorder; cartilage disorder; rheumatoid arthritis; obesity;
KW sports injury; osteoarthritis; hyper-insulinaemia; hypo-insulinaemia;
KW hearing loss; coagulation disorder; stroke; heart attack; cardiant;
KW antidiabetic; anorectic; vulnerary; antiarthritic; osteopathic;
KW antirheumatic; auditory; cerebroprotective; angiogenic.
XX
OS Homo sapiens.
XX
PN US2003004311-A1.
XX
PD 02-JAN-2003.
XX
PF 19-DEC-2001; 2001US-00028072.
XX
PR 18-JUN-1997; 97US-0049911P.
PR 26-AUG-1997; 97US-0056974P.
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 19-SEP-1997; 97US-0059352P.
PR 19-SEP-1997; 97US-0059588P.
PR 24-SEP-1997; 97US-0059836P.
PR 17-OCT-1997; 97US-0062250P.
PR 17-OCT-1997; 97US-0062285P.

PR 17-OCT-1997; 97US-0062287P.
PR 17-OCT-1997; 97US-0063755P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063082P.
PR 24-OCT-1997; 97US-0063127P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063561P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063733P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 11-DEC-1997; 97US-0069212P.
PR 11-DEC-1997; 97US-0069278P.
PR 11-DEC-1997; 97US-0069334P.
PR 16-DEC-1997; 97US-0069694P.
PR 23-JAN-1998; 98US-0072320P.
PR 04-FEB-1998; 98US-0073612P.
PR 09-FEB-1998; 98US-0074086P.
PR 09-FEB-1998; 98US-0074092P.
PR 12-MAR-1998; 98US-0077791P.
PR 20-MAR-1998; 98US-0078910P.
PR 25-MAR-1998; 98US-0079294P.
PR 27-MAR-1998; 98US-0079663P.
PR 27-MAR-1998; 98US-0079728P.
PR 31-MAR-1998; 98US-0080165P.
PR 12-JUN-1998; 98WO-US012456.
PR 14-JUL-1998; 98WO-US014552.
PR 28-AUG-1998; 98WO-US017888.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019093.
PR 14-SEP-1998; 98WO-US019094.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 07-OCT-1998; 98WO-US021141.
PR 29-OCT-1998; 98WO-US022991.
PR 29-OCT-1998; 98WO-US022992.
PR 20-NOV-1998; 98WO-US024855.
PR 01-DEC-1998; 98WO-US025108.
PR 05-JAN-1999; 99WO-US000106.
PR 08-MAR-1999; 99WO-US005028.
PR 10-MAR-1999; 99WO-US005190.
PR 20-APR-1999; 99WO-US008615.
PR 14-MAY-1999; 99WO-US010733.
PR 02-JUN-1999; 99WO-US012252.
PR 01-SEP-1999; 99WO-US020111.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 30-NOV-1999; 99WO-US028409.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 02-DEC-1999; 99WO-US028551.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 30-DEC-1999; 99WO-US031243.
PR 30-DEC-1999; 99WO-US031274.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000277.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 18-FEB-2000; 2000WO-US004342.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 01-MAR-2000; 2000WO-US005601.
PR 02-MAR-2000; 2000WO-US005746.
XX
PA (GETH) GENENTECH INC.
XX
PI Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX
DR WPI; 2003-352836/33.
DR N-PSDB; ACA67170.
XX
PT New isolated PRO polypeptide useful for treating diabetes, rheumatoid
PT arthritis, sports injuries, obesity, hearing loss in mammals, stroke, or
PT heart attack.
XX
PS Claim 12; Fig 354; 643pp; English.
XX
CC The present invention relates to the isolation of novel human PRO
CC polypeptides, and the polynucleotide sequences encoding them. The PRO
CC polypeptides are secreted and transmembrane proteins. The PRO
CC polypeptides and polynucleotides are useful for preparing a medicament
CC useful in the treatment of diabetes, bone and/or cartilage disorders
CC (e.g. rheumatoid arthritis, sports injuries, osteoarthritis), obesity,
CC hyper- or hypo-insulinaemia, hearing loss, and coagulation disorders
CC (e.g. stroke, heart attack). Anti-PRO antibodies are useful in diagnostic
CC assays for PRO, by detecting its expression in specific cells, tissues or
CC serum, and for affinity purification of PRO from recombinant cell culture
CC or natural sources. ABU80870-ABU81144 represent the human PRO
CC polypeptides of the invention. Note: The sequence data for this patent
CC was obtained in electronic format directly from the USPTO web site at
CC seqdata.uspto.gov/psipsDIDEntry.html
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||
Db 493 GVSLSKLSLHN 503

RESULT 23
ABU71906
ID ABU71906 standard; protein; 696 AA.
XX
AC ABU71906;
XX
DT 12-JUN-2003 (first entry)
XX
DE Human secreted/transmembrane protein PRO266.
XX
KW Human; secreted protein; transmembrane protein; PRO; gene therapy;
KW chromosome identification; chromosome marker.
XX
OS Homo sapiens.
XX
PN US2003003530-A1.
XX

PD 02-JAN-2003.
XX
PF 11-JUL-2001; 2001US-00904011.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-329602/31.
DR N-PSDB; ACA60074.
XX
PT New transmembrane polypeptides and nucleic acids encoding the
PT polypeptides, useful in gene therapy, in chromosome identification, as
PT chromosome markers, in generating probes and in tissue typing.
XX
PS Claim 12; Fig 34; 484pp; English.
XX
CC The invention relates to an isolated nucleic acid with at least 80%
CC nucleic acid sequence identity to a nucleotide sequence encoding one of
CC 61 secreted/transmembrane polypeptides, or PRO polypeptides or encoding a
CC PRO protein extracellular domain. Also included are a vector comprising
CC the PRO nucleic acid, a host cell comprising the vector, producing a PRO
CC polypeptide (by culturing the host cell for the expression of the PRO
CC polypeptide, and recovering the PRO polypeptide from the cell culture),
CC an isolated PRO polypeptide (having at least 80% sequence identity to: (
CC a) an amino acid sequence selected from the 61 PRO proteins; (b) an amino
CC acid sequence encoded by a nucleic acid molecule deposited with an ATCC
CC number (detailed in the specification); or (c) an extracellular domain of
CC a PRO polypeptide or to a PRO polypeptide lacking its associated signal
CC peptide), a chimaeric molecule comprising a PRO polypeptide of fused to a
CC heterologous amino acid sequence, an anti-PRO antibody, detecting a
CC PRO245 or PRO1868 in a sample suspected of containing the polypeptide,
CC linking a bioactive molecule to a cell expressing a PRO245 or PRO1868 and
CC modulating at least one biological activity of a cell expressing a PRO245
CC or PRO1868. Nucleic acids which encode PRO can be used to generate either
CC transgenic animals or knock-out animals which may be used in the
CC development and screening of therapeutically useful reagents. The nucleic
CC acids may also be used in gene therapy, in chromosome identification, as
CC chromosome markers, or in generating probes. The PRO polypeptides are
CC useful as molecular markers for protein electrophoresis, and the isolated
CC nucleic acids may be used for recombinantly expressing those markers. The
CC PRO polypeptides and nucleic acids may also be used in tissue typing.
CC Anti-PRO antibodies are useful in diagnostic assays for PRO, and in
CC affinity purification of PRO from recombinant cell culture or natural
CC sources. The present sequence represents a PRO protein
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||
Db 493 GVSLSKLSLHN 503

RESULT 24
ABO01789
ID ABO01789 standard; protein; 696 AA.
XX
AC ABO01789;

XX
DT 07-AUG-2003 (first entry)
XX
DE Novel human secreted and transmembrane protein PRO266.
XX
KW Human; secreted and transmembrane protein; PRO; pharmaceutical;
KW diagnostic; biosensor; bioreactor; Parkinson's disease;
KW Alzheimer's disease; inflammation; nephritis; wound healing;
KW nerve repair; collateral blood vessel formation; cancer;
KW colorectal cancer; haemorrhage; rheumatoid arthritis; diabetes;
KW cirrhosis; fibrosis; restenosis; dermal fibrotic condition; keloid;
KW scarring; ischaemia; stroke; hypertension; heart attack; atherosclerosis;
KW infertility; gene therapy.
XX
OS Homo sapiens.
XX
PN US2002197671-A1.
XX
PD 26-DEC-2002.
XX
PF 17-JUL-2001; 2001US-00907824.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-370793/35.
DR N-PSDB; ACD07474.
XX
PT New genes and secreted and transmembrane polypeptides (e.g. PRO245 or
PT PRO335), useful for treating or diagnosing e.g. Alzheimer's disease,
PT cancers, hemorrhage, rheumatoid arthritis, diabetes, cirrhosis, ischemia
PT or strokes.
XX
PS Claim 12; Fig 34; 482pp; English.
XX
CC The invention describes a new isolated nucleic acid molecule comprising
CC the full length coding sequence of the DNA deposited with the American
CC Type Culture Collection (e.g. ATCC Deposit No. 209258) ,or a sequence
CC with at least 80% identity to a DNA encoding a PRO polypeptide comprising
CC any of 61 sequences having 164-1119 amino acids fully defined in the
CC specification. The PRO polypeptides or polynucleotides are useful as
CC pharmaceuticals, diagnostics, biosensors or bioreactors. These are
CC particularly useful for detecting or treating e.g. Parkinson's disease,
CC Alzheimer's disease, inflammations, nephritis, wound healing, nerve
CC repair, collateral blood vessel formation, cancers (e.g. colorectal
CC cancer), haemorrhage (or reduce risk for haemorrhage), rheumatoid
CC arthritis, diabetes, cirrhosis of the liver, fibrosis of the lungs,
CC restenosis, dermal fibrotic conditions (e.g. keloids or scarring),
CC ischaemia, strokes, hypertension, heart attacks, atherosclerosis, or
CC infertility in mammals (e.g. humans, dogs, cats, cattle, horses, sheep,
CC pigs, goats, or rabbits) The PRO polypeptides are useful as targets for
CC therapeutic intervention in these diseases, and diagnostic determination
CC of the presence of these diseases. The PRO polypeptides are also useful
CC as molecular weight markers, or for chromosome identification. The PRO
CC genes are useful as hybridisation probes, or for screening libraries of
CC human cDNA, genomic DNA or mRNA. The PRO genes may also be used in gene
CC therapy, particularly for replacing a defective gene. This is the amino
CC acid sequence of a novel human secreted and transmembrane PRO polypeptide
XX
SQ Sequence 696 AA;

```
  Query Match             56.8%;  Score 42;  DB 6;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 25
ABU66746
ID   ABU66746 standard; protein; 696 AA.
XX
AC   ABU66746;
XX
DT   23-MAY-2003  (first entry)
XX
DE   Human PRO polypeptide #177.
XX
KW   Human; PRO polypeptide; secreted and transmembrane protein;
KW   tumour necrosis factor-alpha; TNF-alpha; blood; proliferation;
KW   differentiation; chondrocyte; tumour; genetic disorder; cytostatic.
XX
OS   Homo sapiens.
XX
PN   US2003036180-A1.
XX
PD   20-FEB-2003.
XX
PF   09-MAY-2002; 2002US-00143114.
XX
PR   31-MAR-1997;   97WO-US005230.
PR   12-JUN-1998;   98WO-US012456.
PR   14-JUL-1998;   98WO-US014552.
PR   28-AUG-1998;   98WO-US017888.
PR   10-SEP-1998;   98WO-US018824.
PR   14-SEP-1998;   98WO-US019093.
PR   14-SEP-1998;   98WO-US019094.
PR   14-SEP-1998;   98WO-US019177.
PR   16-SEP-1998;   98WO-US019330.
PR   17-SEP-1998;   98WO-US019437.
PR   07-OCT-1998;   98WO-US021141.
PR   29-OCT-1998;   98WO-US022991.
PR   29-OCT-1998;   98WO-US022992.
PR   20-NOV-1998;   98WO-US024855.
PR   01-DEC-1998;   98WO-US025108.
PR   05-JAN-1999;   99WO-US000106.
PR   08-MAR-1999;   99WO-US005028.
PR   10-MAR-1999;   99WO-US005190.
PR   20-APR-1999;   99WO-US008615.
PR   14-MAY-1999;   99WO-US010733.
PR   02-JUN-1999;   99WO-US012252.
PR   01-SEP-1999;   99WO-US020111.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   30-NOV-1999;   99WO-US028409.
PR   01-DEC-1999;   99WO-US028301.
PR   01-DEC-1999;   99WO-US028634.
PR   02-DEC-1999;   99WO-US028551.
PR   02-DEC-1999;   99WO-US028564.
PR   02-DEC-1999;   99WO-US028565.
PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   22-DEC-1999;   99WO-US030720.
PR   30-DEC-1999;   99WO-US031243.
PR   30-DEC-1999;   99WO-US031274.
PR   05-JAN-2000; 2000WO-US000219.
PR   06-JAN-2000; 2000WO-US000277.
PR   06-JAN-2000; 2000WO-US000376.
PR   11-FEB-2000; 2000WO-US003565.
PR   18-FEB-2000; 2000WO-US004341.
PR   18-FEB-2000; 2000WO-US004342.
PR   22-FEB-2000; 2000WO-US004414.
PR   24-FEB-2000; 2000WO-US004914.
PR   24-FEB-2000; 2000WO-US005004.
PR   01-MAR-2000; 2000WO-US005601.
PR   02-MAR-2000; 2000WO-US005746.
PR   02-MAR-2000; 2000WO-US005841.
PR   10-MAR-2000; 2000WO-US006319.
PR   15-MAR-2000; 2000WO-US006884.
PR   20-MAR-2000; 2000WO-US007377.
PR   21-MAR-2000; 2000WO-US007532.
PR   30-MAR-2000; 2000WO-US008439.
PR   17-MAY-2000; 2000WO-US013705.
PR   22-MAY-2000; 2000WO-US014042.
PR   30-MAY-2000; 2000WO-US014941.
PR   02-JUN-2000; 2000WO-US015264.
PR   28-JUL-2000; 2000WO-US020710.
PR   11-AUG-2000; 2000WO-US022031.
PR   23-AUG-2000; 2000WO-US023522.
PR   24-AUG-2000; 2000WO-US023328.
PR   08-NOV-2000; 2000WO-US030952.
PR   10-NOV-2000; 2000WO-US030873.
PR   01-DEC-2000; 2000WO-US032678.
PR   20-DEC-2000; 2000US-00747259.
PR   20-DEC-2000; 2000WO-US034956.
PR   28-FEB-2001; 2001US-00796498.
PR   28-FEB-2001; 2001WO-US006520.
PR   01-MAR-2001; 2001WO-US006666.
PR   09-MAR-2001; 2001US-00802706.
PR   14-MAR-2001; 2001US-00808689.
PR   22-MAR-2001; 2001US-00816744.
PR   05-APR-2001; 2001US-00828366.
PR   10-MAY-2001; 2001US-00854208.
PR   10-MAY-2001; 2001US-00854280.
PR   18-MAY-2001; 2001US-00860216.
PR   25-MAY-2001; 2001US-00866028.
PR   25-MAY-2001; 2001US-00866034.
PR   25-MAY-2001; 2001WO-US017092.
PR   01-JUN-2001; 2001US-00872035.
PR   01-JUN-2001; 2001WO-US017800.
PR   05-JUN-2001; 2001US-00874503.
PR   14-JUN-2001; 2001US-00882636.
PR   19-JUN-2001; 2001US-00886342.
PR   20-JUN-2001; 2001WO-US019692.
PR   21-JUN-2001; 2001US-00887879.
PR   22-JUN-2001; 2001WO-US020116.
PR   29-JUN-2001; 2001WO-US021066.
PR   09-JUL-2001; 2001WO-US021735.
PR   18-JUL-2001; 2001US-00908827.
PR   06-AUG-2001; 2001US-00924419.
PR   09-AUG-2001; 2001US-00927796.
PR   16-AUG-2001; 2001US-00931836.
PR   19-DEC-2001; 2001US-00028072.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Baker KP,  Beresini M,  Deforge L,  Desnoyers L,  Filvaroff E,  Gao W;
PI   Gerritsen ME,  Goddard A,  Godowski PJ,  Gurney AL,  Sherwood S;
PI   Smith V,  Stewart TA,  Tumas D,  Watanabe CK,  Wood WI,  Zhang Z;
XX
DR   WPI; 2003-332040/31.
DR   N-PSDB; ACA03779.
XX
PT   New secreted and transmembrane PRO nucleic acids, useful for gene
PT   therapy, in chromosome and gene mapping, as chromosome markers, in tissue
PT   typing, and in chromosome identification.
XX
PS   Claim 12; Fig 354; 660pp; English.
XX
```

CC The present invention relates to the isolation of novel human PRO
CC polypeptides, and the polynucleotide sequences encoding them. The PRO
CC polypeptides are secreted and transmembrane proteins. The PRO
CC polypeptides are useful for detecting other PRO polypeptides, for linking
CC bioactive molecules to cells expressing PRO polypeptides, for modulating
CC biological activities of cells expressing PRO polypeptides, and for for
CC identifying agonists or antagonists. The PRO polypeptides are useful for
CC for stimulating the release of tumour necrosis factor (TNF)-alpha from
CC human blood, for stimulating the proliferation or differentiation of
CC chondrocytes, and detecting the presence of tumours. The polynucleotide
CC sequences encoding PRO polypeptides are useful as hybridisation probes,
CC in chromosome and gene mapping, in the generation of antisense RNA and
CC DNA, in the preparation of PRO polypeptides, for generating transgenic
CC animals or knockout animals, for the genetic analysis of individuals with
CC genetic disorders, and in gene therapy. ABU66570-ABU66844 represent the
CC human PRO polypeptides of the invention. Note: The sequence data for this
CC patent was obtained in electronic format directly from the USPTO web site
CC at seqdata.uspto.gov/psipsDIDEntry.html
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||
Db 493 GVSLSKLSLHN 503

RESULT 26
ABU54362
ID ABU54362 standard; protein; 696 AA.
XX
AC ABU54362;
XX
DT 10-MAR-2003 (first entry)
XX
DE Human secreted/transmembrane protein PRO266.
XX
KW Human; PRO; secreted protein; transmembrane protein; enterocolitis;
KW gastrointestinal ulceration; skin disease;
KW abnormal keratinocyte differentiation; psoriasis; epithelial cancer;
KW squamous cell carcinoma; Alzheimer's disease; Parkinson's disease;
KW amyotrophic lateral sclerosis; inflammatory disease;
KW rheumatoid arthritis; asthma; multiple sclerosis; organ failure;
KW atherosclerosis; cardiac injury; infertility; birth defect;
KW premature aging; AIDS; acquired immunodeficiency syndrome; cancer;
KW diabetic complication; wound repair.
XX
OS Homo sapiens.
XX
PN US2002132240-A1.
XX
PD 19-SEP-2002.
XX
PF 18-JUL-2001; 2001US-00909320.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 06-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX

DR WPI; 2003-147434/14.
DR N-PSDB; ABX71522.
XX
PT New PRO polypeptides and nucleic acid molecules, useful in diagnosing or
PT treating inflammatory diseases, organ failure, atherosclerosis, cardiac
PT injury, infertility, cancer, AIDS, Alzheimer's disease or Parkinson's
PT disease.
XX
PS Claim 12; Fig 34; 473pp; English.
XX
CC The invention relates to an isolated PRO polypeptide having at least 80%
CC amino acid sequence identity to: (a) any one of 61 fully defined amino
CC acid sequences given in the specification (appearing as ABU54347-
CC ABU54407); (b) an amino acid sequence encoded by the nucleotide sequence
CC deposited under American Type Culture Collection (accession numbers
CC listed in the specification); (c) any one of the PRO sequences which
CC lacks its associated signal peptide; (d) an extracellular domain of the
CC PRO polypeptide with its associated signal peptide; or (e) an
CC extracellular domain of the PRO polypeptide which lacks its associated
CC signal peptide. Also include are the nucleic acids encoding the PRO
CC polypeptides, vectors, host cells and anti-PRO antibodies. The PRO
CC polypeptides and nucleic acids are useful in diagnosing or treating
CC enterocolitis, gastrointestinal ulceration, skin diseases associated with
CC abnormal keratinocyte differentiation, e.g. psoriasis or epithelial
CC cancers such as squamous cell carcinoma, Alzheimer's disease, Parkinson's
CC disease, amyotrophic lateral sclerosis, inflammatory diseases, e.g.
CC rheumatoid arthritis, asthma or multiple sclerosis, organ failure,
CC atherosclerosis, cardiac injury, infertility, birth defects, premature
CC aging, AIDS, cancer, diabetic complications, or mutations in general. The
CC polypeptides are also useful for wound repair and associated therapies
CC concerned with re-growth of tissue. The nucleotide sequences may be used
CC as hybridisation probes in chromosome and gene mapping, or in generating
CC antisense RNA and DNA. PRO nucleic acids are also useful in preparing PRO
CC polypeptides, in assays to identify other proteins or molecules involved
CC in binding reaction, to generate transgenic animals or knockout animals,
CC which in turn are useful in the development and screening of
CC therapeutically useful reagents, for chromosome identification, and
CC tissue typing. The PRO polypeptides and nucleic acid molecules are also
CC useful in gene therapy, and as molecular weight markers for protein
CC electrophoresis purposes. The anti-PRO antibodies may be used in
CC diagnostic assays for PRO, or for the affinity purification of PRO from
CC recombinant cell culture or natural sources. The present sequence
CC represents a PRO polypeptide
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 27
ABO47377
ID ABO47377 standard; protein; 696 AA.
XX
AC ABO47377;
XX
DT 08-OCT-2003 (first entry)
XX
DE Human secreted/transmembrane polypeptide PRO266.
XX
KW Human; abnormal bleeding; gynaecological disease; asthma; hysterectomy;
KW angiogenesis; coronary ischaemic condition; skin disease;
KW gastrointestinal mucosa disorder; acute mucosal lesion; neuropathy; ALS;
KW chronic mucosal lesion; abnormal keratinocyte differentiation; psoriasis;
KW Parkinson's disease; Alzheimer's disease; amyotrophic lateral sclerosis;
KW uncontrolled cell growth; cancer; blood coagulation cascade; thrombosis;
KW haemorrhage; endometrial bleeding; angiogenesis; wound healing; tumour;
KW tissue repair; rheumatoid arthritis; multiple sclerosis; tissue typing.
XX
OS Homo sapiens.
XX
PN US2003044839-A1.
XX
PD 06-MAR-2003.
XX
PF 10-JUL-2001; 2001US-00902903.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.

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PR   26-JUL-1999;   99US-0145698P.
PR   28-JUL-1999;   99US-0146222P.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   01-DEC-1999;   99WO-US028301.
PR   02-DEC-1999;   99WO-US028564.
PR   02-DEC-1999;   99WO-US028565.
PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   05-JAN-2000; 2000WO-US000219.
PR   11-FEB-2000; 2000WO-US003565.
PR   22-FEB-2000; 2000WO-US004414.
PR   24-FEB-2000; 2000WO-US005004.
PR   02-MAR-2000; 2000WO-US005841.
PR   20-MAR-2000; 2000WO-US007377.
PR   30-MAR-2000; 2000WO-US008439.
PR   22-MAY-2000; 2000WO-US014042.
PR   02-JUN-2000; 2000WO-US015264.
PR   28-JUL-2000; 2000WO-US020710.
PR   24-AUG-2000; 2000WO-US023328.
PR   18-SEP-2000; 2000US-00665350.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Ashkenazi A,  Botstein D,  Desnoyers L,  Eaton DL,  Ferrara N;
PI   Filvaroff E,  Fong S,  Gao W,  Gerber H,  Gerritsen ME,  Goddard A;
PI   Godowski PJ,  Grimaldi JC,  Gurney AL,  Hillan KJ,  Kljavin IJ;
PI   Mather JP,  Pan J,  Paoni NF,  Roy MA,  Stewart TA,  Tumas D;
PI   Williams PM,  Wood WI;
XX
DR   WPI; 2003-492258/46.
DR   N-PSDB; ACH06854.
XX
PT   Novel secreted and transmembrane polypeptides and polynucleotides
PT   encoding them useful for treating abnormal bleeding involved in
PT   gynecological diseases, skin diseases and neurodegenerative diseases.
XX
PS   Claim 12; Fig 34; 478pp; English.
XX
CC   The invention relates to an isolated PRO polypeptide. PRO317 is useful in
CC   diagnosing or treating abnormal bleeding involved in gynecological
CC   diseases e.g. to avoid or lessen the need for hysterectomy. PRO317 may
CC   also be useful as an agent that affects angiogenesis and PRO317 is useful
CC   in anti-tumour indications or in treating coronary ischaemic conditions.
CC   PRO211 and PRO217 polypeptides are useful for treating disorders
CC   associated with the preservation and maintenance of gastrointestinal
CC   mucosa and the repair of acute and chronic mucosal lesions, skin diseases
CC   associated with abnormal keratinocyte differentiation (e.g. psoriasis).
CC   PRO187 polypeptide is useful for treating Parkinson's disease,
CC   Alzheimer's disease, amyotrophic lateral sclerosis (ALS), neuropathies
CC   and disease related to uncontrolled cell growth, e.g. cancer. PRO219
CC   polypeptide plays a regulatory role in the blood coagulation cascade.
CC   PRO246 polypeptides which serves as tumour specific antigens may be
CC   exploited as therapeutic targets for anti-tumour drugs. PRO269
CC   polypeptide is useful as an antithrombotic agent with reduced risk for
CC   haemorrhage as compared with heparin. PRO317 polypeptide is useful in
CC   treating endometrial bleeding angiogenesis. PRO287 polypeptides and
CC   portion have therapeutic applications in wound healing and tissue repair.
CC   PRO234 polypeptides are useful for treating asthma, rheumatoid arthritis,
CC   psoriasis and multiple sclerosis. The polypeptide and its nucleic acid
CC   are useful for tissue typing. PRO antibodies are useful for
CC   immunohistochemical staining and/or assay of sample fluids. Anti-PRO
CC   antibodies are useful in diagnostic assays for PRO e.g. detecting its
CC   expression in specific cells, tissues or serum and for affinity
CC   purification of PRO from recombinant cell culture or natural sources. The
CC   present sequence represents the amino acid sequence of a human secreted/
CC   transmembrane PRO polypeptide
XX
SQ   Sequence 696 AA;

  Query Match             56.8%;  Score 42;  DB 6;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 28
ABU59827
ID   ABU59827 standard; protein; 696 AA.
XX
AC   ABU59827;
XX
DT   13-MAY-2003  (first entry)
XX
DE   Novel secreted and transmembrane protein PRO266.
XX
KW   Human; PRO; hypertrophy of neonatal heart; angiogenesis; wound healing;
KW   cardiac insufficiency disorder; cancer; tumour; immune response;
KW   adrenal cortical capillary endothelial growth; c-fos induction;
KW   vascular endothelial growth factor inhibition; VEGF inhibition;
KW   endothelial cell growth inhibitor; T-lymphocytes stimulation;
KW   retinal neurons cell survival; rod photoreceptor cell survival;
KW   retinal disorder; retinitis pigmentosum; kidney disorder;
KW   mammalian kidney mesangial cell proliferation; Berger disease;
KW   dermatitis; herpetiformis; Crohn's disease; chondrocyte proliferation;
KW   chondrocyte redifferentiation; sports injury; arthritis.
XX
OS   Homo sapiens.
XX
PN   US2003017563-A1.
XX
PD   23-JAN-2003.
XX
PF   07-MAY-2002; 2002US-00140808.
XX
PR   31-MAR-1997;   97WO-US005230.
PR   12-JUN-1998;   98WO-US012456.
PR   14-JUL-1998;   98WO-US014552.
PR   28-AUG-1998;   98WO-US017888.
PR   10-SEP-1998;   98WO-US018824.
PR   14-SEP-1998;   98WO-US019093.
PR   14-SEP-1998;   98WO-US019094.
PR   14-SEP-1998;   98WO-US019177.
PR   16-SEP-1998;   98WO-US019330.
PR   17-SEP-1998;   98WO-US019437.
PR   07-OCT-1998;   98WO-US021141.
PR   29-OCT-1998;   98WO-US022991.
PR   29-OCT-1998;   98WO-US022992.
PR   20-NOV-1998;   98WO-US024855.
PR   01-DEC-1998;   98WO-US025108.
PR   05-JAN-1999;   99WO-US000106.
PR   08-MAR-1999;   99WO-US005028.
PR   10-MAR-1999;   99WO-US005190.
PR   20-APR-1999;   99WO-US008615.
PR   14-MAY-1999;   99WO-US010733.
PR   02-JUN-1999;   99WO-US012252.
PR   01-SEP-1999;   99WO-US020111.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   30-NOV-1999;   99WO-US028409.
PR   01-DEC-1999;   99WO-US028301.
PR   01-DEC-1999;   99WO-US028634.
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PR 02-DEC-1999; 99WO-US028551.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 22-DEC-1999; 99WO-US030720.
PR 30-DEC-1999; 99WO-US031243.
PR 30-DEC-1999; 99WO-US031274.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000277.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 18-FEB-2000; 2000WO-US004342.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 01-MAR-2000; 2000WO-US005601.
PR 02-MAR-2000; 2000WO-US005746.
PR 02-MAR-2000; 2000WO-US005841.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
PR 21-MAR-2000; 2000WO-US007532.
PR 30-MAR-2000; 2000WO-US008439.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 10-NOV-2000; 2000WO-US030873.
PR 01-DEC-2000; 2000WO-US032678.
PR 20-DEC-2000; 2000US-00747259.
PR 20-DEC-2000; 2000WO-US034956.
PR 28-FEB-2001; 2001US-00796498.
PR 28-FEB-2001; 2001WO-US006520.
PR 01-MAR-2001; 2001WO-US006666.
PR 09-MAR-2001; 2001US-00802706.
PR 14-MAR-2001; 2001US-00808689.
PR 22-MAR-2001; 2001US-00816744.
PR 05-APR-2001; 2001US-00828366.
PR 10-MAY-2001; 2001US-00854208.
PR 10-MAY-2001; 2001US-00854280.
PR 18-MAY-2001; 2001US-00860216.
PR 25-MAY-2001; 2001US-00866028.
PR 25-MAY-2001; 2001US-00866034.
PR 25-MAY-2001; 2001WO-US017092.
PR 01-JUN-2001; 2001US-00872035.
PR 01-JUN-2001; 2001WO-US017800.
PR 05-JUN-2001; 2001US-00874503.
PR 14-JUN-2001; 2001US-00882636.
PR 19-JUN-2001; 2001US-00886342.
PR 20-JUN-2001; 2001WO-US019692.
PR 21-JUN-2001; 2001US-00887879.
PR 22-JUN-2001; 2001WO-US020116.
PR 29-JUN-2001; 2001WO-US021066.
PR 09-JUL-2001; 2001WO-US021735.
PR 18-JUL-2001; 2001US-00908827.
PR 06-AUG-2001; 2001US-00924419.
PR 09-AUG-2001; 2001US-00927796.
PR 16-AUG-2001; 2001US-00931836.
PR 19-DEC-2001; 2001US-00028072.
XX
PA (GETH) GENENTECH INC.
XX
PI Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX
DR WPI; 2003-148238/14.
DR N-PSDB; ABX89317.
XX
PT Two hundred and seventy five nucleic acids encoding PRO polypeptides,
PT useful for treating pericyte-associated tumors, diabetes and various bone
PT and/or cartilage disorders, e.g. arthritis.
XX
PS Claim 12; Fig 354; 659pp; English.
XX
CC The invention describes an isolated human PRO polypeptide. The PRO
CC polypeptides are useful in detecting PRO polypeptides in a sample, in
CC linking a bioactive molecule to a cell expressing a PRO polypeptide, and
CC in modulating at least one biological activity of a cell expressing a PRO
CC polypeptide. PRO1312 stimulates hypertrophy of neonatal heart and is thus
CC useful for treating cardiac insufficiency disorders. PRO1154 and PRO1186
CC stimulate adrenal cortical capillary endothelial growth, and PRO536,
CC PRO943, PRO828, PRO826, PRO1068 or PRO535, PRO826, PRO819, PRO1126,
CC PRO1360 and PRO1387 induce c-fos in endothelial cells, and are thus
CC useful for treating conditions or disorders where angiogenesis would be
CC beneficial, e.g. wound healing and antagonist of this polypeptide are
CC useful for treating cancerous tumours. PRO812 inhibits vascular
CC endothelial growth factor (VEGF) stimulated proliferation of endothelial
CC cells and is thus useful for inhibiting endothelial cell growth in
CC mammals which would be beneficial in inhibiting tumour growth. PRO826,
CC PRO1068, PRO1184, PRO1346 and PRO1375 stimulate proliferation of
CC stimulated T-lymphocytes and are therapeutically useful for enhancing
CC immune response. PRO828, PRO826, PRO1068 or PRO1132 enhance survival of
CC retinal neurons cells (PRO1132 is also enhances survival/proliferation of
CC rod photoreceptor cells) and therefore are useful for treating retinal
CC disorders of injuries, e.g. retinitis pigmentosum, AMD. PRO819, PRO813
CC and PRO11066 induce proliferation of mammalian kidney mesangial cells,
CC and therefore are useful for treating kidney disorders associated with
CC decreased mesangial cell function such as Berger disease or other
CC nephropathies associated with dermatitis, herpetiformis or Crohn's
CC disease. PRO1310, PRO844, PRO1312, PRO1192 and PRO1387 induce the
CC proliferation and/or redifferentiation of chondrocytes in culture and are
CC thus useful for treating sports injuries, and arthritis. This is the
CC amino acid sequence of a novel human PRO protein
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 29
ABO25017
ID ABO25017 standard; protein; 696 AA.
XX
AC ABO25017;
XX
DT 05-SEP-2003 (first entry)
XX
DE Human secreted/transmembrane protein (PRO) #177.
XX
KW Human; PRO; secreted protein; transmembrane protein; tumour; cytostatic;
KW gene therapy; tumour necrosis factor-alpha; TNF-alpha; blood;
KW proteoglycan; cartilage; cytokine; peripheral blood mononuclear cell;
KW PBMC; glucose uptake; FFA; skeletal muscle cell; adipocyte cell;
KW chondrocyte cell proliferation; chondrocyte cell differentiation;
KW pericyte cell; inner ear utricular supporting cell; T-lymphocyte cell;
KW endothelial cell; A-peptide; factor VIIA.
XX
OS Homo sapiens.
XX
PN US2003036179-A1.
XX

PD 20-FEB-2003.
XX
PF 10-MAY-2002; 2002US-00142431.
XX
PR 31-MAR-1997; 97WO-US005230.
PR 12-JUN-1998; 98WO-US012456.
PR 14-JUL-1998; 98WO-US014552.
PR 28-AUG-1998; 98WO-US017888.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019093.
PR 14-SEP-1998; 98WO-US019094.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 07-OCT-1998; 98WO-US021141.
PR 29-OCT-1998; 98WO-US022991.
PR 29-OCT-1998; 98WO-US022992.
PR 20-NOV-1998; 98WO-US024855.
PR 01-DEC-1998; 98WO-US025108.
PR 05-JAN-1999; 99WO-US000106.
PR 08-MAR-1999; 99WO-US005028.
PR 10-MAR-1999; 99WO-US005190.
PR 20-APR-1999; 99WO-US008615.
PR 14-MAY-1999; 99WO-US010733.
PR 02-JUN-1999; 99WO-US012252.
PR 01-SEP-1999; 99WO-US020111.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 30-NOV-1999; 99WO-US028409.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 02-DEC-1999; 99WO-US028551.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 22-DEC-1999; 99WO-US030720.
PR 30-DEC-1999; 99WO-US031243.
PR 30-DEC-1999; 99WO-US031274.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000277.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 18-FEB-2000; 2000WO-US004342.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 01-MAR-2000; 2000WO-US005601.
PR 02-MAR-2000; 2000WO-US005746.
PR 02-MAR-2000; 2000WO-US005841.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
PR 21-MAR-2000; 2000WO-US007532.
PR 30-MAR-2000; 2000WO-US008439.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 10-NOV-2000; 2000WO-US030873.
PR 01-DEC-2000; 2000WO-US032678.
PR 20-DEC-2000; 2000US-00747259.
PR 20-DEC-2000; 2000WO-US034956.
PR 28-FEB-2001; 2001US-00796498.
PR 28-FEB-2001; 2001WO-US006520.
PR 01-MAR-2001; 2001WO-US006666.
PR 09-MAR-2001; 2001US-00802706.
PR 14-MAR-2001; 2001US-00808689.
PR 22-MAR-2001; 2001US-00816744.
PR 05-APR-2001; 2001US-00828366.
PR 10-MAY-2001; 2001US-00854208.
PR 10-MAY-2001; 2001US-00854280.
PR 18-MAY-2001; 2001US-00860216.
PR 25-MAY-2001; 2001US-00866028.
PR 25-MAY-2001; 2001US-00866034.
PR 25-MAY-2001; 2001WO-US017092.
PR 01-JUN-2001; 2001US-00872035.
PR 01-JUN-2001; 2001WO-US017800.
PR 05-JUN-2001; 2001US-00874503.
PR 14-JUN-2001; 2001US-00882636.
PR 19-JUN-2001; 2001US-00886342.
PR 20-JUN-2001; 2001WO-US019692.
PR 21-JUN-2001; 2001US-00887879.
PR 22-JUN-2001; 2001WO-US020116.
PR 29-JUN-2001; 2001WO-US021066.
PR 09-JUL-2001; 2001WO-US021735.
PR 18-JUL-2001; 2001US-00908827.
PR 06-AUG-2001; 2001US-00924419.
PR 09-AUG-2001; 2001US-00927796.
PR 16-AUG-2001; 2001US-00931836.
PR 19-DEC-2001; 2001US-00028072.
XX
PA (GETH) GENENTECH INC.
XX
PI Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX
DR WPI; 2003-466355/44.
DR N-PSDB; ACD41971.
XX
PT New isolated nucleic acid encoding a PRO polypeptide, e.g. PRO1114 or
PT PRO4978, useful in molecular biology, chromosome and gene mapping, in
PT generating antisense RNA and DNA, and in gene therapy.
XX
PS Claim 12; Fig 354; 659pp; English.
XX
CC The invention relates to an isolated nucleic acid comprising at least 80%
CC sequence identity to a PRO (secreted and transmembrane protein) cDNA
CC comprising a nucleic acid (a) encoding a PRO polypeptide, or its
CC extracellular domain (with or without its associated signal peptide),
CC which comprises any of the 275 120-850 residue amino acid sequences,
CC given in the specification; (b) comprising any of the 275 300-3500
CC nucleotide sequences, given in the specification; or (c) comprising the
CC full-length coding sequence of the nucleotide sequences given in the
CC specification, or of the DNA deposited under any of the American Type
CC Culture Collection (ATCC) Accession Numbers listed in the specification.
CC Also included are a vector comprising the novel nucleic acid, a host cell
CC comprising the vector, producing a PRO polypeptide, the isolated PRO
CC polypeptides detailed above, a chimaeric molecule comprising the PRO
CC polypeptide of fused to a heterologous amino acid sequence, an anti-PRO
CC antibody, detecting a PRO polypeptide in a sample suspected of containing
CC the PRO polypeptide, linking a bioactive molecule to a cell expressing a
CC PRO polypeptide, modulating at least one biological activity of a cell
CC expressing a PRO polypeptide, stimulating the release of tumour necrosis
CC factor-alpha (TNF-alpha) from human blood, (or proteoglycans from
CC cartilage or cytokine from peripheral blood mononuclear cells (PBMC)),
CC modulating the uptake of glucose or FFA by skeletal muscle cells or
CC adipocyte cells, stimulating the proliferation or differentiation of
CC chondrocyte cells (or proliferation of or gene expression in pericyte
CC cells), stimulating the proliferation of inner ear utricular supporting
CC cells (or of T-lymphocyte cells, or of endothelial cells), inhibiting the
CC binding of A-peptide to factor VIIA, or differentiation of adipocyte
CC cells, detecting the presence of a tumour in a mammal and an

CC oligonucleotide probe derived from any of the nucleotide sequences given
CC in the specification. The polynucleotide is useful in molecular biology,
CC including uses as hybridisation probes, in chromosome and gene mapping,
CC in generating antisense RNA and DNA, and in gene therapy. The
CC polynucleotide may also be used in preparing PRO polypeptides by
CC recombinant techniques, and in generating either transgenic animals or
CC knock-out animals which, in turn, are useful in the development and
CC screening of therapeutically useful reagents. The PRO polypeptide or the
CC antibody is used in preparing a medicament for treating a condition
CC responsive to the polypeptide or antibody, such as tumours, and in
CC various diagnostic assays. The present sequence rperesents a PRO
CC polypeptide
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 30
ABU64514
ID ABU64514 standard; protein; 696 AA.
XX
AC ABU64514;
XX
DT 13-MAY-2003 (first entry)
XX
DE Human secreted/transmembrane protein, #18.
XX
KW Human; PRO; secreted; transmembrane; pharmaceutical; diagnostic;
KW biosensor; bioreactor; therapeutic; hyperplasia; endometriosis; cancer;
KW tumour; ischaemia; coronary arterial disease; polycystic kidney disease;
KW renal failure; inflammatory response; asthma; rheumatoid arthritis;
KW psoriasis; multiple sclerosis; gene therapy; cytostatic; gynecological;
KW cardiant; nephrotropic; hepatotropic; antiinflammatory.
XX
OS Homo sapiens.
XX
PN US2002160374-A1.
XX
PD 31-OCT-2002.
XX
PF 12-JUL-2001; 2001US-00905291.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 01-DEC-1998; 98WO-US025108.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-288105/28.
DR N-PSDB; ABX96091.
XX
PT New secreted and transmembrane PRO polypeptides (e.g. PRO533 or PRO245)
PT and genes encoding them, useful for detecting or treating e.g.
PT hyperplasia, endometriosis, cancers, ischemia, coronary arterial disease
PT or inflammations.
XX
PS Claim 12; Fig 34; 477pp; English.

XX
CC The invention discloses isolated PRO secreted/transmembrane polypeptides
CC and the nucleic acid encoding them. The polypeptides can be used to raise
CC antibodies that specifically bind to the PRO polypeptide, for linking a
CC bioactive molecule to a cell expressing a PRO protein and for modulating
CC at least one biological activity of a cell. The PRO polypeptides or
CC polynucleotides are also useful as pharmaceuticals, diagnostics,
CC biosensors or bioreactors, for detecting or treating e.g. hyperplasia,
CC endometriosis, cancers (e.g. those involving solid tumours), ischaemia,
CC coronary arterial disease, polycystic kidney disease, chronic or acute
CC renal failure, or inflammatory responses (e.g. asthma, rheumatoid
CC arthritis, psoriasis or multiple sclerosis) in mammals. The PRO genes may
CC also be used in gene therapy, particularly for replacing a defective
CC gene. The sequences presented in ABU64499-ABU64559 are the PRO
CC polynucleotides of the invention
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||
Db 493 GVSLSKLSLHN 503

RESULT 31
ABU67360
ID ABU67360 standard; protein; 696 AA.
XX
AC ABU67360;
XX
DT 29-MAY-2003 (first entry)
XX
DE Human secreted protein PRO226.
XX
KW Human; gene therapy; mucosal lesion; ulcer; enterocolitis; skin disease;
KW psoriasis; cancer; lung cancer; colon cancer; nerve cell disease;
KW Alzheimer's disease; Parkinson's disease; Usher syndrome; angiogenesis;
KW atrophia areata; inflammatory disease; asthma; rheumatoid arthritis;
KW ischaemia.
XX
OS Homo sapiens.
XX
PN US2003023054-A1.
XX
PD 30-JAN-2003.
XX
PF 16-JUL-2001; 2001US-00906742.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;

```
PI   Filvaroff E,  Fong S,  Gao W,  Gerber H,  Gerritsen ME,  Goddard A;
PI   Godowski PJ,  Grimaldi JC,  Gurney AL,  Hillan KJ,  Kljavin IJ;
PI   Mather JP,  Pan J,  Paoni NF,  Roy MA,  Stewart TA,  Tumas D;
PI   Williams PM,  Wood WI;
XX
DR   WPI; 2003-331485/31.
DR   N-PSDB; ACA05412.
XX
PT   Sixty one isolated nucleic acids encoding a PRO polypeptide, e.g. PRO245
PT   or PRO1868, useful in chromosome and gene mapping, in generating
PT   antisense RNA and DNA, and in treating cancer and Alzheimer's disease.
XX
PS   Example 14; Fig 34; 481pp; English.
XX
CC   The invention relates to sixty one nucleic acids encoding PRO
CC   polypeptides (secreted and transmembrane). The polynucleotide is useful
CC   in molecular biology, including uses as hybridisation probes, in
CC   chromosome and gene mapping, in generating antisense RNA and DNA, and in
CC   gene therapy. The polynucleotide may also be used in preparing PRO
CC   polypeptides by recombinant techniques, and in generating either
CC   transgenic animals or knock-out animals which, in turn, are useful in the
CC   development and screening of therapeutically useful reagents. The PRO
CC   polypeptide or the antibody is used in preparing a medicament for
CC   treating a condition responsive to the polypeptide or antibody, such as
CC   mucosal lesions e.g. ulcers and enterocolitis, skin disease e.g.
CC   psoriasis, cancer e.g. lung cancer and colon cancer, nerve cell disease
CC   e.g. Alzheimer's disease and Parkinson's disease, Usher syndrome,
CC   atrophia areata, angiogenesis, inflammatory disease e.g asthma and
CC   rheumatoid arthritis, ischaemia, and in various diagnostic assays. The
CC   present sequence represents the amino acid sequence of a PRO polypeptide
XX
SQ   Sequence 696 AA;

  Query Match             56.8%;  Score 42;  DB 6;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 32
ABO14880
ID   ABO14880 standard; protein; 696 AA.
XX
AC   ABO14880;
XX
DT   25-AUG-2003  (first entry)
XX
DE   Human secreted / transmembrane polypeptide PRO266.
XX
KW   Human; gene therapy; tumour; tissue typing; obesity; arthritis; diabetes;
KW   hypoinsulinaemia; hyperinsulinaemia; vascular permeability;
KW   cardiac insufficiency disorder; immune response; regeneration; cartilage;
KW   auditory hair cell; hearing loss; bone disorder; sports injury.
XX
OS   Homo sapiens.
XX
PN   US2003036060-A1.
XX
PD   20-FEB-2003.
XX
PF   12-JUL-2001; 2001US-00904859.
XX
PR   17-SEP-1997;   97US-0059113P.
PR   17-SEP-1997;   97US-0059115P.
PR   17-SEP-1997;   97US-0059117P.
PR   17-SEP-1997;   97US-0059119P.
PR   17-SEP-1997;   97US-0059121P.
PR   17-SEP-1997;   97US-0059122P.
PR   17-SEP-1997;   97US-0059184P.
PR   18-SEP-1997;   97US-0059263P.
PR   18-SEP-1997;   97US-0059266P.
PR   15-OCT-1997;   97US-0062125P.
PR   17-OCT-1997;   97US-0062285P.
PR   17-OCT-1997;   97US-0062287P.
PR   21-OCT-1997;   97US-0063486P.
PR   24-OCT-1997;   97US-0062814P.
PR   24-OCT-1997;   97US-0062816P.
PR   24-OCT-1997;   97US-0063045P.
PR   24-OCT-1997;   97US-0063120P.
PR   24-OCT-1997;   97US-0063121P.
PR   24-OCT-1997;   97US-0063127P.
PR   24-OCT-1997;   97US-0063128P.
PR   27-OCT-1997;   97US-0063327P.
PR   27-OCT-1997;   97US-0063329P.
PR   28-OCT-1997;   97US-0063541P.
PR   28-OCT-1997;   97US-0063542P.
PR   28-OCT-1997;   97US-0063544P.
PR   28-OCT-1997;   97US-0063549P.
PR   28-OCT-1997;   97US-0063550P.
PR   28-OCT-1997;   97US-0063564P.
PR   29-OCT-1997;   97US-0063435P.
PR   29-OCT-1997;   97US-0063704P.
PR   29-OCT-1997;   97US-0063732P.
PR   29-OCT-1997;   97US-0063734P.
PR   29-OCT-1997;   97US-0063735P.
PR   29-OCT-1997;   97US-0063738P.
PR   29-OCT-1997;   97US-0064215P.
PR   31-OCT-1997;   97US-0063870P.
PR   31-OCT-1997;   97US-0064103P.
PR   03-NOV-1997;   97US-0064248P.
PR   07-NOV-1997;   97US-0064809P.
PR   12-NOV-1997;   97US-0065186P.
PR   17-NOV-1997;   97US-0065846P.
PR   18-NOV-1997;   97US-0065693P.
PR   21-NOV-1997;   97US-0066120P.
PR   21-NOV-1997;   97US-0066364P.
PR   24-NOV-1997;   97US-0066453P.
PR   24-NOV-1997;   97US-0066466P.
PR   24-NOV-1997;   97US-0066511P.
PR   24-NOV-1997;   97US-0066770P.
PR   24-NOV-1997;   97US-0066772P.
PR   25-NOV-1997;   97US-0066840P.
PR   12-DEC-1997;   97US-0069425P.
PR   04-JUN-1998;   98US-0088026P.
PR   10-SEP-1998;   98US-0099803P.
PR   10-SEP-1998;   98WO-US018824.
PR   14-SEP-1998;   98US-0100262P.
PR   14-SEP-1998;   98WO-US019177.
PR   16-SEP-1998;   98WO-US019330.
PR   17-SEP-1998;   98US-0100858P.
PR   17-SEP-1998;   98WO-US019437.
PR   13-OCT-1998;   98US-0104080P.
PR   20-NOV-1998;   98US-0109304P.
PR   01-DEC-1998;   98WO-US025108.
PR   22-DEC-1998;   98US-0113296P.
PR   07-JUL-1999;   99US-0143048P.
PR   26-JUL-1999;   99US-0145698P.
PR   28-JUL-1999;   99US-0146222P.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   01-DEC-1999;   99WO-US028301.
PR   02-DEC-1999;   99WO-US028564.
PR   02-DEC-1999;   99WO-US028565.
PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   05-JAN-2000; 2000WO-US000219.
PR   11-FEB-2000; 2000WO-US003565.
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PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-417923/39.
DR N-PSDB; ACD20079.
XX
PT Novel secreted and transmembrane polypeptide for modulating biological
PT activity of cell expressing the polypeptide, identifying agonists or
PT antagonists of polypeptide, and as molecular weight markers.
XX
PS Claim 12; Fig 34; 469pp; English.
XX
CC The invention relates to an isolated, secreted and transmembrane
CC polypeptide, termed PRO polypeptide. The polypeptide is useful for
CC identifying agonists or antagonists of the polypeptide, for preparing
CC variants of the polypeptide, as molecular weight markers for protein
CC electrophoresis purpose and the nucleic acid is useful for recombinantly
CC expressing those markers. The polypeptide is also useful as therapeutic
CC agent. PRO is useful in assays to identify other proteins or molecules
CC involved in binding interaction. The nucleic acid is useful as
CC hybridisation probes, in chromosome and gene mapping, in generation of
CC antisense RNA and DNA, in the preparation of PRO polypeptide, for
CC generating transgenic animals or knockout animals which in turn are
CC useful in the development and screening of therapeutically useful
CC reagents, to construct hybridisation probes for mapping the gene which
CC encodes the PRO and for the genetic analysis of individuals with genetic
CC disorders, in gene therapy, for chromosome identification, as chromosome
CC marker, and for generating probes for polymerase chain reaction (PCR),
CC Northern analysis, Southern analysis and Western analysis. PRO antibody
CC is useful in diagnostic assays for PRO, e.g. detecting its expression in
CC specific cells, tissues or serum and for affinity purification of PRO
CC from recombinant cell culture or natural sources. The polypeptide or its
CC antibody is useful for the preparation of medicament for treating
CC conditions which is responsive to the PRO polypeptide or anti-PRO
CC antibody e.g. tumour. The polypeptide and the nucleic acid is useful for
CC tissue typing. The polypeptide is useful for treating obesity, diabetes
CC or hypo- or hyper-insulinaemia and cardiac insufficiency disorders, for
CC inhibiting tumour growth, enhances vascular permeability and immune
CC response, for inducing regeneration of auditory hair cells and for
CC treating hearing loss in mammals and for treating bone and/or cartilage
CC disorders such as sports injuries and arthritis. The present sequence
CC represents the amino acid sequence of a human secreted and transmembrane
CC PRO polypeptide
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||
Db 493 GVSLSKLSLHN 503

RESULT 33
ABU67022

ID ABU67022 standard; protein; 696 AA.
XX
AC ABU67022;
XX
DT 27-MAY-2003 (first entry)
XX
DE Human secreted/transmembrane, PRO, protein SEQ ID 354.
XX
KW Human; secreted protein; transmembrane protein; PRO;
KW inflammatory disease; organ failure; atherosclerosis; cardiac injury;
KW infertility; birth defects; premature aging; AIDS; biosensor;
KW acquired immunodeficiency syndrome; cancer; diabetic complication;
KW bioreactor; tumour.
XX
OS Homo sapiens.
XX
PN US2003032155-A1.
XX
PD 13-FEB-2003.
XX
PF 03-MAY-2002; 2002US-00137865.
XX
PR 31-MAR-1997; 97WO-US005230.
PR 12-JUN-1998; 98WO-US012456.
PR 14-JUL-1998; 98WO-US014552.
PR 28-AUG-1998; 98WO-US017888.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019093.
PR 14-SEP-1998; 98WO-US019094.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 07-OCT-1998; 98WO-US021141.
PR 29-OCT-1998; 98WO-US022991.
PR 29-OCT-1998; 98WO-US022992.
PR 20-NOV-1998; 98WO-US024855.
PR 01-DEC-1998; 98WO-US025108.
PR 05-JAN-1999; 99WO-US000106.
PR 08-MAR-1999; 99WO-US005028.
PR 10-MAR-1999; 99WO-US005190.
PR 20-APR-1999; 99WO-US008615.
PR 14-MAY-1999; 99WO-US010733.
PR 02-JUN-1999; 99WO-US012252.
PR 01-SEP-1999; 99WO-US020111.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 30-NOV-1999; 99WO-US028409.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 02-DEC-1999; 99WO-US028551.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 22-DEC-1999; 99WO-US030720.
PR 30-DEC-1999; 99WO-US031243.
PR 30-DEC-1999; 99WO-US031274.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000277.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 18-FEB-2000; 2000WO-US004342.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 01-MAR-2000; 2000WO-US005601.

PR 02-MAR-2000; 2000WO-US005746.
PR 02-MAR-2000; 2000WO-US005841.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
PR 21-MAR-2000; 2000WO-US007532.
PR 30-MAR-2000; 2000WO-US008439.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 10-NOV-2000; 2000WO-US030873.
PR 01-DEC-2000; 2000WO-US032678.
PR 20-DEC-2000; 2000US-00747259.
PR 20-DEC-2000; 2000WO-US034956.
PR 28-FEB-2001; 2001US-00796498.
PR 28-FEB-2001; 2001WO-US006520.
PR 01-MAR-2001; 2001WO-US006666.
PR 09-MAR-2001; 2001US-00802706.
PR 14-MAR-2001; 2001US-00808689.
PR 22-MAR-2001; 2001US-00816744.
PR 05-APR-2001; 2001US-00828366.
PR 10-MAY-2001; 2001US-00854208.
PR 10-MAY-2001; 2001US-00854280.
PR 18-MAY-2001; 2001US-00860216.
PR 25-MAY-2001; 2001US-00866028.
PR 25-MAY-2001; 2001US-00866034.
PR 25-MAY-2001; 2001WO-US017092.
PR 01-JUN-2001; 2001US-00872035.
PR 01-JUN-2001; 2001WO-US017800.
PR 05-JUN-2001; 2001US-00874503.
PR 14-JUN-2001; 2001US-00882636.
PR 19-JUN-2001; 2001US-00886342.
PR 20-JUN-2001; 2001WO-US019692.
PR 21-JUN-2001; 2001US-00887879.
PR 22-JUN-2001; 2001WO-US020116.
PR 29-JUN-2001; 2001WO-US021066.
PR 09-JUL-2001; 2001WO-US021735.
PR 18-JUL-2001; 2001US-00908827.
PR 06-AUG-2001; 2001US-00924419.
PR 09-AUG-2001; 2001US-00927796.
PR 16-AUG-2001; 2001US-00931836.
PR 19-DEC-2001; 2001US-00028072.
XX
PA (GETH) GENENTECH INC.
XX
PI Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX
DR WPI; 2003-331925/31.
DR N-PSDB; ACA04200.
XX
PT New secreted and transmembrane nucleic acids and polypeptides, designated
PT as PRO, useful for treating inflammation, organ failure, atherosclerosis,
PT cardiac injury, infertility, birth defects, premature aging, AIDS, or
PT cancer.
XX
PS Claim 12; Fig 354; 659pp; English.
XX
CC The invention relates to an isolated nucleic acid comprising, or which is
CC at least 80% identical to, or the full-length coding sequence of, any of
CC the 275 nucleotide sequences, encoding the corresponding PRO polypeptide
CC (one of 275 secreted or transmembrane proteins). The nucleic acid further
CC comprises the full-length coding sequence of the DNA deposited under
CC American Type Culture Collection (ATCC) accession number in a list given
CC in the specification. Also included are vectors and host cells for
CC producing PRO proteins, PRO fusion proteins, anti-PRO antibodies, PRO
CC extracellular domains and mature sequences, methods of detecting PRO
CC proteins, methods for stimulating the release of TNF-alpha (tumour
CC necrosis factor alpha) from human blood, (and the proliferation of
CC differentiation of chondrocyte cells, the proliferation of, or gene
CC expression in pericyte cells, the release or proteoglycans from
CC cartilage, proliferation of inner ear urticular supporting cells, the
CC proliferation of T-lymphocyte cells, the release of a cytokine from
CC peripheral blood mononuclear cells (PBMC), or the proliferation of
CC endothelial cells), a method for modulating the uptake of glucose or free
CC fatty acid (FFA) by skeletal muscle cells, a method for inhibiting the
CC binding of A-peptide to factor VIIA, or the differentiation of adipocyte
CC cells, a method for detecting the presence of a tumour in a mammal and an
CC oligonucleotide probe derived from any of the nucleotide sequences cited
CC above. The nucleic acids and polypeptides are useful for treating
CC inflammatory diseases, organ failure, atherosclerosis, cardiac injury,
CC infertility, birth defects, premature aging, AIDS (acquired
CC immunodeficiency syndrome), cancer, or diabetic complications. The
CC nucleic acids are useful as hybridisation probes, in chromosome and gene
CC mapping, and in generating antisense RNA or DNA. The polypeptides are
CC useful as pharmaceuticals, diagnostics, biosensors or bioreactors. Both
CC are useful in tissue typing. The present sequence represents a PRO
CC protein of the invention
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

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Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
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RESULT 34
ABU69637
ID ABU69637 standard; protein; 696 AA.
XX
AC ABU69637;
XX
DT 05-JUN-2003 (first entry)
XX
DE Novel human secreted and transmembrane protein PRO266.
XX
KW Human; secreted and transmembrane protein; gene therapy; psoriasis;
KW enterocolitis; gastrointestinal ulceration; skin disease;
KW keratinocyte differentiation; epithelial cancer; Alzheimer's disease;
KW squamous cell carcinoma; Parkinson's disease; inflammatory disease;
KW amyotrophic lateral sclerosis; rheumatoid arthritis; asthma;
KW multiple sclerosis; organ failure; atherosclerosis; cardiac injury;
KW infertility; birth defect; premature aging; AIDS; cancer;
KW diabetic complication; wound repair; tissue re-growth.
XX
OS Homo sapiens.
XX
PN US2003017463-A1.
XX
PD 23-JAN-2003.
XX
PF 11-JUL-2001; 2001US-00903640.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.

PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-341586/32.
DR N-PSDB; ACA54882.
XX
PT New PRO polypeptides and nucleic acid molecules, useful in diagnosing or
PT treating inflammatory diseases, organ failure, atherosclerosis, cardiac
PT injury, infertility, cancer, AIDS, Alzheimer's disease or Parkinson's
PT disease.
XX
PS Claim 12; Fig 34; 473pp; English.
XX
CC The invention describes sixty one nucleic acids encoding PRO polypeptides
CC (secreted and transmembrane). The PRO polypeptides and nucleic acids are
CC useful in diagnosing or treating enterocolitis, gastrointestinal
CC ulceration, skin diseases associated with abnormal keratinocyte
CC differentiation, e.g. psoriasis or epithelial cancers such as squamous
CC cell carcinoma, Alzheimer's disease, Parkinson's disease, amyotrophic
CC lateral sclerosis, inflammatory diseases, e.g. rheumatoid arthritis,
CC asthma or multiple sclerosis, organ failure, atherosclerosis, cardiac
CC injury, infertility, birth defects, premature aging, AIDS, cancer,
CC diabetic complications, or mutations in general. The polypeptides are
CC also useful for wound repair and associated therapies concerned with re-
CC growth of tissue. The PRO polypeptides and nucleic acid molecules are
CC also useful in gene therapy, and as molecular weight markers for protein
CC electrophoresis purposes. The anti-PRO antibodies may be used in
CC diagnostic assays for PRO, or for the affinity purification of PRO from
CC recombinant cell culture or natural sources. This is the amino acid
CC sequence of a novel human PRO polypeptide
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 35
ABO14819
ID ABO14819 standard; protein; 696 AA.
XX
AC ABO14819;
XX
DT 22-AUG-2003 (first entry)
XX
DE Human secreted / transmembrane polypeptide PRO266.
XX
KW Human; ss; gene therapy; apoptosis; bleeding; tumour; ALS;
KW gynaecological disease; hysterectomy; angiogenesis; skin disease; cancer;
KW coronary ischaemic condition; gastrointestinal mucosa disorder; asthma;
KW mucosal lesion repair; keratinocyte differentiation; psoriasis;
KW Parkinson's disease; Alzheimer's disease; amyotrophic lateral sclerosis;
KW neuropathy; blood coagulation cascade disorder; thrombosis; haemorrhage;
KW neurodegenerative disease; endometrial bleeding; wound healing;
KW tissue repair; rheumatoid arthritis; multiple sclerosis; tissue typing.
XX

OS Homo sapiens.
XX
PN US2003027143-A1.
XX
PD 06-FEB-2003.
XX
PF 16-JUL-2001; 2001US-00906838.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
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PR 29-OCT-1997; 97US-0063435P.
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PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.
PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-417249/39.
DR N-PSDB; ACD19717.
XX
PT Novel secreted and transmembrane polypeptides and polynucleotides
PT encoding them useful for treating abnormal bleeding involved in
PT gynecological diseases, skin diseases and neurodegenerative diseases.
XX
PS Claim 12; Fig 34; 467pp; English.
XX
CC The invention relates to an isolated secreted and transmembrane PRO
CC polypeptide. The PRO polypeptides are useful for modulating biological
CC activity of a cell, in diagnosing or treating abnormal bleeding involved
CC in gynaecological diseases e.g. to avoid or lessen the need for
CC hysterectomy, for treating angiogenesis, tumour, coronary ischaemic
CC condition, disorders associated with the preservation and maintenance of
CC gastrointestinal mucosa and the repair of acute and chronic mucosal
CC lesions, skin diseases associated with abnormal keratinocyte
CC differentiation (e.g. psoriasis), Parkinson's disease, Alzheimer's
CC disease, amyotrophic lateral sclerosis (ALS), neuropathies, disease
CC related to uncontrolled cell growth (e.g. cancer), blood coagulation
CC cascade disorders, neurodegenerative disease, thrombosis, haemorrhage,
CC endometrial bleeding, wound healing, tissue repair, asthma, rheumatoid
CC arthritis, multiple sclerosis. Nucleic acid encoding PRO polypeptides are
CC useful in molecular biology including uses as hybridisation probes and in
CC the generation of antisense RNA and DNA, for preparing PRO polypeptides,
CC for generating transgenic animals or knockout animals. The PRO
CC polypeptides and their nucleic acids are useful for tissue typing. PRO
CC antibodies are useful for immunohistochemical staining and/or assay of
CC sample fluids. Anti-PRO antibodies are useful in diagnostic assays for
CC PRO e.g. detecting its expression in specific cells, tissues or serum and
CC for affinity purification of PRO from recombinant cell culture or natural
CC sources. The present sequence represents the amino acid sequence of a
CC human secreted and transmembrane PRO polypeptide
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;

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  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 36
ADA45873
ID   ADA45873 standard; protein; 696 AA.
XX
AC   ADA45873;
XX
DT   20-NOV-2003  (first entry)
XX
DE   Novel human secreted and transmembrane protein PRO266.
XX
KW   Human; secreted and transmembrane protein; PRO;
KW   Tumour necrosis factor alpha release; TNF-alpha release;
KW   glucose uptake modulator; FFA uptake modulator;
KW   cell proliferation stimulator; cell differentiation stimulator;
KW   cell differentiation inihibitor; cytokine release stimulator; tumour;
KW   lung tumoue; colon tumour; breast tumour; prostate tumour; rectal tumour;
KW   cervical tumour; liver tumour; chromosome mapping; gene mapping;
KW   gene therapy; chromosome identification; chromosome marker.
XX
OS   Homo sapiens.
XX
PN   US2003022328-A1.
XX
PD   30-JAN-2003.
XX
PF   16-APR-2002; 2002US-00123904.
XX
PR   31-MAR-1997;   97WO-US005230.
PR   12-JUN-1998;   98WO-US012456.
PR   14-JUL-1998;   98WO-US014552.
PR   28-AUG-1998;   98WO-US017888.
PR   10-SEP-1998;   98WO-US018824.
PR   14-SEP-1998;   98WO-US019093.
PR   14-SEP-1998;   98WO-US019094.
PR   14-SEP-1998;   98WO-US019177.
PR   16-SEP-1998;   98WO-US019330.
PR   17-SEP-1998;   98WO-US019437.
PR   07-OCT-1998;   98WO-US021141.
PR   29-OCT-1998;   98WO-US022991.
PR   29-OCT-1998;   98WO-US022992.
PR   20-NOV-1998;   98WO-US024855.
PR   01-DEC-1998;   98WO-US025108.
PR   05-JAN-1999;   99WO-US000106.
PR   08-MAR-1999;   99WO-US005028.
PR   10-MAR-1999;   99WO-US005190.
PR   20-APR-1999;   99WO-US008615.
PR   14-MAY-1999;   99WO-US010733.
PR   02-JUN-1999;   99WO-US012252.
PR   01-SEP-1999;   99WO-US020111.
PR   08-SEP-1999;   99WO-US020594.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   15-SEP-1999;   99WO-US021547.
PR   05-OCT-1999;   99WO-US023089.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   30-NOV-1999;   99WO-US028409.
PR   01-DEC-1999;   99WO-US028301.
PR   01-DEC-1999;   99WO-US028634.
PR   02-DEC-1999;   99WO-US028551.
PR   02-DEC-1999;   99WO-US028564.
PR   02-DEC-1999;   99WO-US028565.
PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   22-DEC-1999;   99WO-US030720.
PR   30-DEC-1999;   99WO-US031243.
PR   30-DEC-1999;   99WO-US031274.
PR   05-JAN-2000; 2000WO-US000219.
PR   06-JAN-2000; 2000WO-US000277.
PR   06-JAN-2000; 2000WO-US000376.
PR   11-FEB-2000; 2000WO-US003565.
PR   18-FEB-2000; 2000WO-US004341.
PR   18-FEB-2000; 2000WO-US004342.
PR   22-FEB-2000; 2000WO-US004414.
PR   24-FEB-2000; 2000WO-US004914.
PR   24-FEB-2000; 2000WO-US005004.
PR   01-MAR-2000; 2000WO-US005601.
PR   02-MAR-2000; 2000WO-US005746.
PR   02-MAR-2000; 2000WO-US005841.
PR   10-MAR-2000; 2000WO-US006319.
PR   15-MAR-2000; 2000WO-US006884.
PR   20-MAR-2000; 2000WO-US007377.
PR   21-MAR-2000; 2000WO-US007532.
PR   30-MAR-2000; 2000WO-US008439.
PR   17-MAY-2000; 2000WO-US013705.
PR   22-MAY-2000; 2000WO-US014042.
PR   30-MAY-2000; 2000WO-US014941.
PR   02-JUN-2000; 2000WO-US015264.
PR   28-JUL-2000; 2000WO-US020710.
PR   11-AUG-2000; 2000WO-US022031.
PR   23-AUG-2000; 2000WO-US023522.
PR   24-AUG-2000; 2000WO-US023328.
PR   08-NOV-2000; 2000WO-US030952.
PR   10-NOV-2000; 2000WO-US030873.
PR   01-DEC-2000; 2000WO-US032678.
PR   20-DEC-2000; 2000US-00747259.
PR   20-DEC-2000; 2000WO-US034956.
PR   28-FEB-2001; 2001US-00796498.
PR   28-FEB-2001; 2001WO-US006520.
PR   01-MAR-2001; 2001WO-US006666.
PR   09-MAR-2001; 2001US-00802706.
PR   14-MAR-2001; 2001US-00808689.
PR   22-MAR-2001; 2001US-00816744.
PR   05-APR-2001; 2001US-00828366.
PR   10-MAY-2001; 2001US-00854208.
PR   10-MAY-2001; 2001US-00854280.
PR   18-MAY-2001; 2001US-00860216.
PR   25-MAY-2001; 2001US-00866028.
PR   25-MAY-2001; 2001US-00866034.
PR   25-MAY-2001; 2001WO-US017092.
PR   01-JUN-2001; 2001US-00872035.
PR   01-JUN-2001; 2001WO-US017800.
PR   05-JUN-2001; 2001US-00874503.
PR   14-JUN-2001; 2001US-00882636.
PR   19-JUN-2001; 2001US-00886342.
PR   20-JUN-2001; 2001WO-US019692.
PR   21-JUN-2001; 2001US-00887879.
PR   22-JUN-2001; 2001WO-US020116.
PR   29-JUN-2001; 2001WO-US021066.
PR   09-JUL-2001; 2001WO-US021735.
PR   18-JUL-2001; 2001US-00908827.
PR   06-AUG-2001; 2001US-00924419.
PR   09-AUG-2001; 2001US-00927796.
PR   16-AUG-2001; 2001US-00931836.
PR   19-DEC-2001; 2001US-00028072.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Baker KP,  Beresini M,  Deforge L,  Desnoyers L,  Filvaroff E,  Gao W;
PI   Gerritsen ME,  Goddard A,  Godowski PJ,  Gurney AL,  Sherwood S;
PI   Smith V,  Stewart TA,  Tumas D,  Watanabe CK,  Wood WI,  Zhang Z;
XX
DR   WPI; 2003-584997/55.
DR   N-PSDB; ADA45872.
XX
PT   Novel secreted and transmembrane polypeptide for modulating biological
PT   activity of cell expressing the polypeptide, identifying agonists or
PT   antagonists of polypeptide, and as molecular weight markers.
```

XX
PS Claim 12; Fig 354; 659pp; English.
XX
CC The invention describes 305 nucleic acids encoding PRO (secreted and
CC transmembrane) polypeptides (I). (I) is useful for stimulating the
CC release of TNF-alpha from human blood, for modulating the uptake of
CC glucose or FFA by skeletal muscle cells or adipocyte cells, for
CC stimulating the proliferation or differentiation of chondrocyte cells,
CC for stimulating the proliferation of or gene expression in pericyte
CC cells, for stimulating the release of proteoglycans from cartilage, for
CC stimulating the proliferation of inner ear utricular supporting cells,
CC for stimulating the proliferation of T-lymphocyte cells, for stimulating
CC the release of a cytokine from PBMC cells, for inhibiting the binding of
CC A-peptide to factor VIIA, for inhibiting the differentiation of adipocyte
CC cells, for stimulating proliferation of endothelial cells, for detecting
CC the presence of tumour in a mammal. The tumour is lung, colon, breast,
CC prostate, rectal, cervical or liver tumour. The oligonucleotide probes
CC are useful for isolating genomic and cDNA nucleotide sequences or
CC antisense probes. (I) is also useful as therapeutic agent. PRO is useful
CC in assays to identify other proteins or molecules involved in binding
CC interaction. A polynucleotide (II) encoding (I) is useful in chromosome
CC and gene mapping, in generation of antisense RNA and DNA, in the
CC preparation of PRO polypeptide, for generating transgenic animals or
CC knockout animals which in turn are useful in the development and
CC screening of therapeutically useful reagents, in gene therapy, for
CC chromosome identification, as chromosome marker, and for generating
CC probes. An anti-(I)-antibody is useful in diagnostic assays for PRO, e.g.
CC detecting its expression in specific cells, tissues or serum, and for
CC affinity purification of PRO from recombinant cell culture or natural
CC sources. (I) and (II) are useful for tissue typing. This is the amino
CC acid sequence of a novel human secreted and transmembrane PRO
CC polypeptide.
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

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Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
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RESULT 37
ADA76304
ID ADA76304 standard; protein; 696 AA.
XX
AC ADA76304;
XX
DT 20-NOV-2003 (first entry)
XX
DE Human PRO polypeptide #177.
XX
KW Human; PRO; secreted polypeptide; transmembrane polypeptide;
KW tumour necrosis factor-alpha; TNF-alpha; chondrocyte cell; tumour;
KW cancer; adrenal; lung; colon; breast; prostate; rectum; kidney; cervix;
KW liver; microvascular endothelial cell; glucose; FFA;
KW skeletal muscle cell; adipocyte cell; pericyte cell;
KW inner ear utricular supporting cell; T-lymphocyte cell;
KW endothelial cell tube formation; bone disorder; cartilage disorder;
KW sports injury; proteoglycan; articular cartilage defect; osteoarthritis;
KW rheumatoid arthritis; haemoglobin-associated disorder thalassaemia;
KW immune system cell infiltration.
XX
OS Homo sapiens.
XX
PN US2003073212-A1.
XX
PD 17-APR-2003.
XX
PF 16-APR-2002; 2002US-00123903.
XX
PR 31-MAR-1997; 97WO-US005230.
PR 12-JUN-1998; 98WO-US012456.
PR 14-JUL-1998; 98WO-US014552.
PR 28-AUG-1998; 98WO-US017888.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019093.
PR 14-SEP-1998; 98WO-US019094.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 07-OCT-1998; 98WO-US021141.
PR 29-OCT-1998; 98WO-US022991.
PR 29-OCT-1998; 98WO-US022992.
PR 20-NOV-1998; 98WO-US024855.
PR 01-DEC-1998; 98WO-US025108.
PR 05-JAN-1999; 99WO-US000106.
PR 08-MAR-1999; 99WO-US005028.
PR 10-MAR-1999; 99WO-US005190.
PR 20-APR-1999; 99WO-US008615.
PR 14-MAY-1999; 99WO-US010733.
PR 02-JUN-1999; 99WO-US012252.
PR 01-SEP-1999; 99WO-US020111.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 30-NOV-1999; 99WO-US028409.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 02-DEC-1999; 99WO-US028551.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 22-DEC-1999; 99WO-US030720.
PR 30-DEC-1999; 99WO-US031243.
PR 30-DEC-1999; 99WO-US031274.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000277.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 18-FEB-2000; 2000WO-US004342.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 01-MAR-2000; 2000WO-US005601.
PR 02-MAR-2000; 2000WO-US005746.
PR 02-MAR-2000; 2000WO-US005841.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
PR 21-MAR-2000; 2000WO-US007532.
PR 30-MAR-2000; 2000WO-US008439.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 10-NOV-2000; 2000WO-US030873.
PR 01-DEC-2000; 2000WO-US032678.
PR 20-DEC-2000; 2000US-00747259.
PR 20-DEC-2000; 2000WO-US034956.
PR 28-FEB-2001; 2001US-00796498.
PR 28-FEB-2001; 2001WO-US006520.

PR 01-MAR-2001; 2001WO-US006666.
PR 09-MAR-2001; 2001US-00802706.
PR 14-MAR-2001; 2001US-00808689.
PR 22-MAR-2001; 2001US-00816744.
PR 05-APR-2001; 2001US-00828366.
PR 10-MAY-2001; 2001US-00854208.
PR 10-MAY-2001; 2001US-00854280.
PR 18-MAY-2001; 2001US-00860216.
PR 25-MAY-2001; 2001US-00866028.
PR 25-MAY-2001; 2001US-00866034.
PR 25-MAY-2001; 2001WO-US017092.
PR 01-JUN-2001; 2001US-00872035.
PR 01-JUN-2001; 2001WO-US017800.
PR 05-JUN-2001; 2001US-00874503.
PR 14-JUN-2001; 2001US-00882636.
PR 19-JUN-2001; 2001US-00886342.
PR 20-JUN-2001; 2001WO-US019692.
PR 21-JUN-2001; 2001US-00887879.
PR 22-JUN-2001; 2001WO-US020116.
PR 29-JUN-2001; 2001WO-US021066.
PR 09-JUL-2001; 2001WO-US021735.
PR 18-JUL-2001; 2001US-00908827.
PR 06-AUG-2001; 2001US-00924419.
PR 09-AUG-2001; 2001US-00927796.
PR 16-AUG-2001; 2001US-00931836.
PR 19-DEC-2001; 2001US-00028072.
XX
PA (GETH) GENENTECH INC.
XX
PI Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX
DR WPI; 2003-687639/65.
DR N-PSDB; ADA76303.
XX
PT New isolated nucleic acid encoding a secreted and transmembrane
PT polypeptide, designated e.g. PRO1114 or PRO4978, useful in chromosome and
PT gene mapping, in generating antisense RNA and DNA, and in gene therapy.
XX
PS Claim 12; Fig 354; 659pp; English.
XX
CC The invention relates to isolated human PRO polypeptides (secreted and
CC transmembrane polypeptides) and the polynucleotides encoding them. The
CC invention also relates to an antibody which specifically binds to a PRO
CC polypeptide, a method for stimulating the release of tumour necrosis
CC factor-alpha (TNF-alpha) from human blood, a method for stimulating the
CC proliferation or differentiation of chondrocyte cells and a method for
CC detecting the presence of a tumour in a mammal (e.g. adrenal, lung,
CC colon, breast, prostate, rectal, kidney, cervical and liver tumours). The
CC polynucleotides are useful in molecular biology, including uses as
CC hybridisation probes, in chromosome and gene mapping, in generating
CC antisense RNA and DNA and in gene therapy. The polynucleotides may also
CC be used in preparing PRO polypeptides by recombinant techniques and in
CC generating either transgenic animals or knock-out animals which are
CC useful in the development and screening of therapeutically useful
CC reagents. The PRO polypeptides or antibodies are used in preparing a
CC medicament for treating a condition responsive to the polypeptides or
CC antibodies, such as tumours, for stimulating and inhibiting proliferation
CC of human microvascular endothelial cells, for modulating the uptake of
CC glucose or FFA by skeletal muscle cells or adipocyte cells, for
CC stimulating differentiation of adipocyte cells, for stimulating
CC proliferation of or gene expression in pericyte cells, for stimulating
CC the proliferation of inner ear utricular supporting cells or T-lymphocyte
CC cells, for inducing endothelial cell tube formation and for treating
CC various bone and/or cartilage disorders such as sports injuries and
CC arthritis. PRO polypeptides which stimulate the release of proteoglycans
CC from cartilage are useful for treating sports-related joint problems,
CC articular cartilage defects, osteoarthritis and rheumatoid arthritis. PRO
CC polypeptides are also useful for treating various mammalian haemoglobin-
CC associated disorders such as various thalassaemias and conditions which
CC may benefit from enhanced local immune system cell infiltration. This
CC sequence represents a human PRO polypeptide of the invention. Note: The
CC sequence data for this patent is also available in electronic format from
CC USPTO at seqdata.uspto.gov/sequence.html.
XX
SQ Sequence 696 AA;

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  Query Match             56.8%;  Score 42;  DB 6;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
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RESULT 38
ADB29296
ID ADB29296 standard; protein; 696 AA.
XX
AC ADB29296;
XX
DT 20-NOV-2003 (first entry)
XX
DE Human secreted/transmembrane protein, #18.
XX
KW Human; PRO; secreted; transmembrane; gastrointestinal mucosa;
KW mucosal lesion; skin disease; keratinocyte differentiation; psoriasis;
KW Parkinson's disease; Alzheimer's diseases; amyotrophic lateral sclerosis;
KW ALS; neuropathy; cell growth; cancer; tumour; viral infection;
KW neurodegenerative disease; antithrombotic agent; haemorrhage;
KW endometrial bleeding angiogenesis; kidney tissue; apoptosis; therapeutic;
KW tissue typing; immunohistochemical staining; gene therapy; nootropic;
KW neuroprotective; cytostatic; virucide; anticoagulant.
XX
OS Homo sapiens.
XX
PN US2003092002-A1.
XX
PD 15-MAY-2003.
XX
PF 10-JUL-2001; 2001US-00902615.
XX
PR 17-SEP-1997; 97US-0059113P.
PR 17-SEP-1997; 97US-0059115P.
PR 17-SEP-1997; 97US-0059117P.
PR 17-SEP-1997; 97US-0059119P.
PR 17-SEP-1997; 97US-0059121P.
PR 17-SEP-1997; 97US-0059122P.
PR 17-SEP-1997; 97US-0059184P.
PR 18-SEP-1997; 97US-0059263P.
PR 18-SEP-1997; 97US-0059266P.
PR 15-OCT-1997; 97US-0062125P.
PR 17-OCT-1997; 97US-0062285P.
PR 17-OCT-1997; 97US-0062287P.
PR 21-OCT-1997; 97US-0063486P.
PR 24-OCT-1997; 97US-0062814P.
PR 24-OCT-1997; 97US-0062816P.
PR 24-OCT-1997; 97US-0063045P.
PR 24-OCT-1997; 97US-0063120P.
PR 24-OCT-1997; 97US-0063121P.
PR 24-OCT-1997; 97US-0063127P.
PR 24-OCT-1997; 97US-0063128P.
PR 27-OCT-1997; 97US-0063327P.
PR 27-OCT-1997; 97US-0063329P.
PR 28-OCT-1997; 97US-0063541P.
PR 28-OCT-1997; 97US-0063542P.
PR 28-OCT-1997; 97US-0063544P.
PR 28-OCT-1997; 97US-0063549P.
PR 28-OCT-1997; 97US-0063550P.
PR 28-OCT-1997; 97US-0063564P.
PR 29-OCT-1997; 97US-0063435P.
PR 29-OCT-1997; 97US-0063704P.
PR 29-OCT-1997; 97US-0063732P.
PR 29-OCT-1997; 97US-0063734P.

PR 29-OCT-1997; 97US-0063735P.
PR 29-OCT-1997; 97US-0063738P.
PR 29-OCT-1997; 97US-0064215P.
PR 31-OCT-1997; 97US-0063870P.
PR 31-OCT-1997; 97US-0064103P.
PR 03-NOV-1997; 97US-0064248P.
PR 07-NOV-1997; 97US-0064809P.
PR 12-NOV-1997; 97US-0065186P.
PR 17-NOV-1997; 97US-0065846P.
PR 18-NOV-1997; 97US-0065693P.
PR 21-NOV-1997; 97US-0066120P.
PR 21-NOV-1997; 97US-0066364P.
PR 24-NOV-1997; 97US-0066453P.
PR 24-NOV-1997; 97US-0066466P.
PR 24-NOV-1997; 97US-0066511P.
PR 24-NOV-1997; 97US-0066770P.
PR 24-NOV-1997; 97US-0066772P.
PR 25-NOV-1997; 97US-0066840P.
PR 12-DEC-1997; 97US-0069425P.
PR 04-JUN-1998; 98US-0088026P.
PR 10-SEP-1998; 98US-0099803P.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98US-0100262P.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 13-OCT-1998; 98US-0104080P.
PR 20-NOV-1998; 98US-0109304P.
PR 01-DEC-1998; 98WO-US025108.
PR 22-DEC-1998; 98US-0113296P.
PR 07-JUL-1999; 99US-0143048P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 05-JAN-2000; 2000WO-US000219.
PR 11-FEB-2000; 2000WO-US003565.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US005044.
PR 02-MAR-2000; 2000WO-US005841.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 22-MAY-2000; 2000WO-US014042.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 24-AUG-2000; 2000WO-US023328.
PR 18-SEP-2000; 2000US-00665350.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi A, Botstein D, Desnoyers L, Eaton DL, Ferrara N;
PI Filvaroff E, Fong S, Gao W, Gerber H, Gerritsen ME, Goddard A;
PI Godowski PJ, Grimaldi JC, Gurney AL, Hillan KJ, Kljavin IJ;
PI Mather JP, Pan J, Paoni NF, Roy MA, Stewart TA, Tumas D;
PI Williams PM, Wood WI;
XX
DR WPI; 2003-765473/72.
DR N-PSDB; ADB29295.
XX
PT Novel isolated native PRO polypeptide useful for treating Parkinson's
PT disease, enterocolitis, Zollinger-Ellison syndrome gastrointestinal
PT ulceration, Alzheimer's disease, amyotrophic lateral sclerosis, Usher
PT syndrome.
XX
PS Claim 12; Fig 34; 469pp; English.
XX
CC The invention discloses isolated PRO secreted/transmembrane polypeptides
CC and the nucleic acid encoding them. The polypeptides can be used to raise
CC antibodies that specifically bind to the PRO polypeptide, for linking a
CC bioactive molecule to a cell expressing a PRO protein and for modulating
CC at least one biological activity of a cell. PRO polypeptides are useful
CC for detecting other PRO polypeptides in a sample and for linking a
CC bioactive molecule to a cell expressing a PRO polypeptide. The PRO
CC polypeptide antibodies are useful for modulating the biological activity
CC of a cell expressing PRO polypeptides. PRO polypeptides are also useful
CC for treating disorders associated with the preservation and maintenance
CC of gastrointestinal mucosa and the repair of acute and chronic mucosal
CC lesions, skin diseases associated with abnormal keratinocyte
CC differentiation (e.g. psoriasis), Parkinson's disease, Alzheimer's
CC diseases, amyotrophic lateral sclerosis (ALS), neuropathies and
CC additionally, disease related to uncontrolled cell growth, e.g. cancer.
CC PRO polypeptides also serves as tumour specific antigens which may be
CC exploited as therapeutic targets for anti-tumour drugs, and are also
CC employed therapeutically in vivo for lessening the effects of viral
CC infection. The PRO polypeptides can be also used in assays to determine
CC if it has a role in neurodegenerative diseases or their reversal, as an
CC antithrombotic agent with reduced risk for haemorrhage as compared with
CC heparin, in treating other PRO-associated disorders, in modulating
CC endometrial bleeding angiogenesis, and may also have an effect on kidney
CC tissue. PRO polypeptides and their portions affect the expression of
CC genes which have a role in apoptosis. The polynucleotides are useful in
CC molecular biology including uses as hybridisation probes for cDNA library
CC to isolate the full-length PRO cDNA or to isolate other cDNAs, in
CC chromosome and gene mapping, in the generation of antisense RNA and DNA,
CC for preparing PRO polypeptides, for generating transgenic animals or
CC knockout animals which are useful in the development and screening of
CC therapeutically useful reagents, as probes and for the genetic analysis
CC of individuals with genetic disorders as well as for recombinantly
CC expressing the protein and for chromosome identification. The proteins
CC are useful as molecular marker for protein electrophoresis purposes, as
CC therapeutic agents, for screening compounds to identify those that mimic
CC the PRO polypeptide (agonists) or prevent the effect of the PRO
CC polypeptide (antagonists). The polynucleotides and proteins are useful
CC for tissue typing. PRO antibodies are useful for immunohistochemical
CC staining and/or assay of sample fluids. Anti-PRO antibodies are useful in
CC diagnostic assays for PRO e.g. detecting its expression in specific
CC cells, tissues or serum and for affinity purification of PRO from
CC recombinant cell culture or natural sources. The PRO genes may also be
CC used in gene therapy, particularly for replacing a defective gene. The
CC sequence presented is a PRO polynucleotide of the invention.
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

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Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
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RESULT 39
ADA18954
ID ADA18954 standard; protein; 696 AA.
XX
AC ADA18954;
XX
DT 20-NOV-2003 (first entry)
XX
DE Human PRO polypeptide #177.
XX
KW Human; PRO; secreted polypeptide; transmembrane polypeptide;
KW tumour necrosis factor-alpha; TNF-alpha; blood; chondrocyte cell; lung;

KW colon; breast; prostate; rectum; cervix; liver; tumour; cancer;
KW glucose uptake; FFA; adipocyte cell; pericyte cell; proteoglycan;
KW cartilage; inner ear utricular supporting cell; cytokine; A-peptide;
KW factor VIIA; endothelial cell.
XX
OS Homo sapiens.
XX
PN US2003054517-A1.
XX
PD 20-MAR-2003.
XX
PF 08-MAY-2002; 2002US-00141755.
XX
PR 31-MAR-1997; 97WO-US005230.
PR 12-JUN-1998; 98WO-US012456.
PR 14-JUL-1998; 98WO-US014552.
PR 28-AUG-1998; 98WO-US017888.
PR 10-SEP-1998; 98WO-US018824.
PR 14-SEP-1998; 98WO-US019093.
PR 14-SEP-1998; 98WO-US019094.
PR 14-SEP-1998; 98WO-US019177.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 07-OCT-1998; 98WO-US021141.
PR 29-OCT-1998; 98WO-US022991.
PR 29-OCT-1998; 98WO-US022992.
PR 20-NOV-1998; 98WO-US024855.
PR 01-DEC-1998; 98WO-US025108.
PR 05-JAN-1999; 99WO-US000106.
PR 08-MAR-1999; 99WO-US005028.
PR 10-MAR-1999; 99WO-US005190.
PR 20-APR-1999; 99WO-US008615.
PR 14-MAY-1999; 99WO-US010733.
PR 02-JUN-1999; 99WO-US012252.
PR 01-SEP-1999; 99WO-US020111.
PR 08-SEP-1999; 99WO-US020594.
PR 13-SEP-1999; 99WO-US020944.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 05-OCT-1999; 99WO-US023089.
PR 29-NOV-1999; 99WO-US028214.
PR 30-NOV-1999; 99WO-US028313.
PR 30-NOV-1999; 99WO-US028409.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 02-DEC-1999; 99WO-US028551.
PR 02-DEC-1999; 99WO-US028564.
PR 02-DEC-1999; 99WO-US028565.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 20-DEC-1999; 99WO-US030999.
PR 22-DEC-1999; 99WO-US030720.
PR 30-DEC-1999; 99WO-US031243.
PR 30-DEC-1999; 99WO-US031274.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000277.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 18-FEB-2000; 2000WO-US004342.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 01-MAR-2000; 2000WO-US005601.
PR 02-MAR-2000; 2000WO-US005746.
PR 02-MAR-2000; 2000WO-US005841.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
PR 21-MAR-2000; 2000WO-US007532.
PR 30-MAR-2000; 2000WO-US008439.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 10-NOV-2000; 2000WO-US030873.
PR 01-DEC-2000; 2000WO-US032678.
PR 20-DEC-2000; 2000US-00747259.
PR 20-DEC-2000; 2000WO-US034956.
PR 28-FEB-2001; 2001US-00796498.
PR 28-FEB-2001; 2001WO-US006520.
PR 01-MAR-2001; 2001WO-US006666.
PR 09-MAR-2001; 2001US-00802706.
PR 14-MAR-2001; 2001US-00808689.
PR 22-MAR-2001; 2001US-00816744.
PR 05-APR-2001; 2001US-00828366.
PR 10-MAY-2001; 2001US-00854208.
PR 10-MAY-2001; 2001US-00854280.
PR 18-MAY-2001; 2001US-00860216.
PR 25-MAY-2001; 2001US-00866028.
PR 25-MAY-2001; 2001US-00866034.
PR 25-MAY-2001; 2001WO-US017092.
PR 01-JUN-2001; 2001US-00872035.
PR 01-JUN-2001; 2001WO-US017800.
PR 05-JUN-2001; 2001US-00874503.
PR 14-JUN-2001; 2001US-00882636.
PR 19-JUN-2001; 2001US-00886342.
PR 20-JUN-2001; 2001WO-US019692.
PR 21-JUN-2001; 2001US-00887879.
PR 22-JUN-2001; 2001WO-US020116.
PR 29-JUN-2001; 2001WO-US021066.
PR 09-JUL-2001; 2001WO-US021735.
PR 18-JUL-2001; 2001US-00908827.
PR 06-AUG-2001; 2001US-00924419.
PR 09-AUG-2001; 2001US-00927796.
PR 16-AUG-2001; 2001US-00931836.
PR 19-DEC-2001; 2001US-00028072.
XX
PA (GETH) GENENTECH INC.
XX
PI Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX
DR WPI; 2003-521854/49.
DR N-PSDB; ADA18953.
XX
PT New PRO nucleic acid, useful for preparing a composition for treating
PT e.g., tumors.
XX
PS Claim 12; Fig 354; 660pp; English.
XX
CC The invention relates to isolated human PRO polypeptides (secreted and
CC transmembrane polypeptides) and the polynucleotides encoding them. The
CC invention also relates to an antibody which specifically binds to a PRO
CC polypeptide, a method for stimulating the release of tumour necrosis
CC factor-alpha (TNF-alpha) from human blood, a method for stimulating the
CC proliferation or differentiation of chondrocyte cells and a method for
CC detecting the presence of a tumour in a mammal (e.g. lung, colon, breast,
CC prostate, rectal, cervical and liver tumours). The polynucleotides are
CC useful in molecular biology, including uses as hybridisation probes, in
CC chromosome and gene mapping, in generating antisense RNA and DNA and in
CC gene therapy. The polynucleotides may also be used in preparing PRO
CC polypeptides by recombinant techniques and in generating either
CC transgenic animals or knock-out animals which are useful in the
CC development and screening of therapeutically useful reagents. The PRO
CC polypeptides or antibodies are used in preparing a medicament for
CC treating a condition responsive to the polypeptides or antibodies, such
CC as tumours, for modulating the uptake of glucose or FFA by adipocyte
CC cells, for stimulating the proliferation of or gene expression in
CC pericyte cells, for stimulating the release of proteoglycans from

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CC   cartilage, for stimulating the proliferation of inner ear utricular
CC   supporting cells, for stimulating the release of cytokines from PBMC
CC   cells, for inhibiting the binding of A-peptide to factor VIIA, for
CC   inhibiting the differentiation of adipocyte cells and for stimulating the
CC   proliferation of endothelial cells. This sequence represents a human PRO
CC   polypeptide of the invention. Note: The sequence data for this patent is
CC   also available in electronic format from USPTO at
CC   seqdata.uspto.gov/sequence.html.
XX
SQ   Sequence 696 AA;

  Query Match             56.8%;  Score 42;  DB 6;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 67;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
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Db        493 GVSLSKLSLHN 503


RESULT 40
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ID   ADA61577 standard; protein; 696 AA.
XX
AC   ADA61577;
XX
DT   20-NOV-2003  (first entry)
XX
DE   Homo sapiens.
XX
KW   Human; secreted and transmembrane protein; PRO;
KW   Tumour necrosis factor alpha release; TNF-alpha release;
KW   glucose uptake modulator; FFA uptake modulator;
KW   cell proliferation stimulator; cell differentiation stimulator;
KW   cell differentiation inihibitor; cytokine release stimulator; tumour;
KW   lung tumoue; colon tumour; breast tumour; prostate tumour; rectal tumour;
KW   cervical tumour; liver tumour; chromosome mapping; gene mapping;
KW   gene therapy; chromosome identification; chromosome marker.
XX
OS   Novel.
OS   human.
OS   secreted.
OS   and.
OS   transmembrane.
OS   protein.
OS   PRO266.
XX
PN   US2003049816-A1.
XX
PD   13-MAR-2003.
XX
PF   15-APR-2002; 2002US-00123262.
XX
PR   31-MAR-1997;   97WO-US005230.
PR   12-JUN-1998;   98WO-US012456.
PR   14-JUL-1998;   98WO-US014552.
PR   28-AUG-1998;   98WO-US017888.
PR   10-SEP-1998;   98WO-US018824.
PR   14-SEP-1998;   98WO-US019093.
PR   14-SEP-1998;   98WO-US019094.
PR   14-SEP-1998;   98WO-US019177.
PR   16-SEP-1998;   98WO-US019330.
PR   17-SEP-1998;   98WO-US019437.
PR   07-OCT-1998;   98WO-US021141.
PR   29-OCT-1998;   98WO-US022991.
PR   29-OCT-1998;   98WO-US022992.
PR   20-NOV-1998;   98WO-US024855.
PR   01-DEC-1998;   98WO-US025108.
PR   05-JAN-1999;   99WO-US000106.
PR   08-MAR-1999;   99WO-US005028.
PR   10-MAR-1999;   99WO-US005190.
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PR   14-MAY-1999;   99WO-US010733.
PR   02-JUN-1999;   99WO-US012252.
PR   01-SEP-1999;   99WO-US020111.
PR   08-SEP-1999;   99WO-US020594.
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PR   05-OCT-1999;   99WO-US023089.
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PR   02-DEC-1999;   99WO-US028551.
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PR   16-DEC-1999;   99WO-US030095.
PR   20-DEC-1999;   99WO-US030911.
PR   20-DEC-1999;   99WO-US030999.
PR   22-DEC-1999;   99WO-US030720.
PR   30-DEC-1999;   99WO-US031243.
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PR   05-JAN-2000; 2000WO-US000219.
PR   06-JAN-2000; 2000WO-US000277.
PR   06-JAN-2000; 2000WO-US000376.
PR   11-FEB-2000; 2000WO-US003565.
PR   18-FEB-2000; 2000WO-US004341.
PR   18-FEB-2000; 2000WO-US004342.
PR   22-FEB-2000; 2000WO-US004414.
PR   24-FEB-2000; 2000WO-US004914.
PR   24-FEB-2000; 2000WO-US005004.
PR   01-MAR-2000; 2000WO-US005601.
PR   02-MAR-2000; 2000WO-US005746.
PR   02-MAR-2000; 2000WO-US005841.
PR   10-MAR-2000; 2000WO-US006319.
PR   15-MAR-2000; 2000WO-US006884.
PR   20-MAR-2000; 2000WO-US007377.
PR   21-MAR-2000; 2000WO-US007532.
PR   30-MAR-2000; 2000WO-US008439.
PR   17-MAY-2000; 2000WO-US013705.
PR   22-MAY-2000; 2000WO-US014042.
PR   30-MAY-2000; 2000WO-US014941.
PR   02-JUN-2000; 2000WO-US015264.
PR   28-JUL-2000; 2000WO-US020710.
PR   11-AUG-2000; 2000WO-US022031.
PR   23-AUG-2000; 2000WO-US023522.
PR   24-AUG-2000; 2000WO-US023328.
PR   08-NOV-2000; 2000WO-US030952.
PR   10-NOV-2000; 2000WO-US030873.
PR   01-DEC-2000; 2000WO-US032678.
PR   20-DEC-2000; 2000US-00747259.
PR   20-DEC-2000; 2000WO-US034956.
PR   28-FEB-2001; 2001US-00796498.
PR   28-FEB-2001; 2001WO-US006520.
PR   01-MAR-2001; 2001WO-US006666.
PR   09-MAR-2001; 2001US-00802706.
PR   14-MAR-2001; 2001US-00808689.
PR   22-MAR-2001; 2001US-00816744.
PR   05-APR-2001; 2001US-00828366.
PR   10-MAY-2001; 2001US-00854208.
PR   10-MAY-2001; 2001US-00854280.
PR   18-MAY-2001; 2001US-00860216.
PR   25-MAY-2001; 2001US-00866028.
PR   25-MAY-2001; 2001US-00866034.
PR   25-MAY-2001; 2001WO-US017092.
PR   01-JUN-2001; 2001US-00872035.
PR   01-JUN-2001; 2001WO-US017800.
PR   05-JUN-2001; 2001US-00874503.
PR   14-JUN-2001; 2001US-00882636.
PR   19-JUN-2001; 2001US-00886342.
PR   20-JUN-2001; 2001WO-US019692.
PR   21-JUN-2001; 2001US-00887879.
PR   22-JUN-2001; 2001WO-US020116.
PR   29-JUN-2001; 2001WO-US021066.
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PR 09-JUL-2001; 2001WO-US021735.
PR 18-JUL-2001; 2001US-00908827.
PR 06-AUG-2001; 2001US-00924419.
PR 09-AUG-2001; 2001US-00927796.
PR 16-AUG-2001; 2001US-00931836.
PR 19-DEC-2001; 2001US-00028072.
XX
PA (GETH) GENENTECH INC.
XX
PI Baker KP, Beresini M, Deforge L, Desnoyers L, Filvaroff E, Gao W;
PI Gerritsen ME, Goddard A, Godowski PJ, Gurney AL, Sherwood S;
PI Smith V, Stewart TA, Tumas D, Watanabe CK, Wood WI, Zhang Z;
XX
DR WPI; 2003-695892/66.
DR N-PSDB; ADA61576.
XX
PT New PRO nucleic acid and encode polypeptides, are useful for
PT manufacturing a medicament for diagnosing or treating cancer.
XX
PS Claim 12; Fig 354; 660pp; English.
XX
CC The invention describes 305 nucleic acids encoding PRO (secreted and
CC transmembrane) polypeptides (I). (I) is useful for stimulating the
CC release of TNF-alpha from human blood, for modulating the uptake of
CC glucose or FFA by skeletal muscle cells or adipocyte cells, for
CC stimulating the proliferation or differentiation of chondrocyte cells,
CC for stimulating the proliferation of or gene expression in pericyte
CC cells, for stimulating the release of proteoglycans from cartilage, for
CC stimulating the proliferation of inner ear utricular supporting cells,
CC for stimulating the proliferation of T-lymphocyte cells, for stimulating
CC the release of a cytokine from PBMC cells, for inhibiting the binding of
CC A-peptide to factor VIIA, for inhibiting the differentiation of adipocyte
CC cells, for stimulating proliferation of endothelial cells, for detecting
CC the presence of tumour in a mammal. The tumour is lung, colon, breast,
CC prostate, rectal, cervical or liver tumour. The oligonucleotide probes
CC are useful for isolating genomic and cDNA nucleotide sequences or
CC antisense probes. (I) is also useful as therapeutic agent. PRO is useful
CC in assays to identify other proteins or molecules involved in binding
CC interaction. A polynucleotide (II) encoding (I) is useful in chromosome
CC and gene mapping, in generation of antisense RNA and DNA, in the
CC preparation of PRO polypeptide, for generating transgenic animals or
CC knockout animals which in turn are useful in the development and
CC screening of therapeutically useful reagents, in gene therapy, for
CC chromosome identification, as chromosome marker, and for generating
CC probes. An anti-(I)-antibody is useful in diagnostic assays for PRO, e.g.
CC detecting its expression in specific cells, tissues or serum, and for
CC affinity purification of PRO from recombinant cell culture or natural
CC sources. (I) and (II) are useful for tissue typing. This is the amino
CC acid sequence of a novel human secreted and transmembrane PRO
CC polypeptide.
XX
SQ Sequence 696 AA;

Query Match 56.8%; Score 42; DB 6; Length 696;
Best Local Similarity 63.6%; Pred. No. 67;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

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Qy          4 GMALSKINLHN 14
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Db        493 GVSLSKLSLHN 503
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OM protein - protein search, using sw model

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Title: US-10-769-514-17
Perfect score: 74
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Searched: 1867569 seqs, 417829326 residues

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Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 100 summaries

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2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
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6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	74	100.0	14	5	US-10-769-514-17	Sequence 17, Appl
2	74	100.0	15	5	US-10-769-514-15	Sequence 15, Appl
3	74	100.0	15	5	US-10-769-514-16	Sequence 16, Appl
4	44	59.5	64	4	US-10-425-115-367739	Sequence 367739,
5	43	58.1	71	4	US-10-424-599-250435	Sequence 250435,
6	42	56.8	635	4	US-10-424-233-19	Sequence 19, Appl
7	42	56.8	695	3	US-09-374-046A-132	Sequence 132, App
8	42	56.8	695	4	US-10-616-263-132	Sequence 132, App
9	42	56.8	696	3	US-09-909-320-91	Sequence 91, Appl
10	42	56.8	696	3	US-09-909-088B-91	Sequence 91, Appl
11	42	56.8	696	3	US-09-905-291A-91	Sequence 91, Appl
12	42	56.8	696	3	US-09-902-853-91	Sequence 91, Appl
13	42	56.8	696	3	US-09-907-824-91	Sequence 91, Appl
14	42	56.8	696	3	US-09-907-841-91	Sequence 91, Appl
15	42	56.8	696	3	US-09-904-011-91	Sequence 91, Appl
16	42	56.8	696	3	US-09-903-640-91	Sequence 91, Appl
17	42	56.8	696	3	US-09-908-093-91	Sequence 91, Appl
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19	42	56.8	696	3	US-09-906-838-91	Sequence 91, Appl
20	42	56.8	696	3	US-09-907-613-91	Sequence 91, Appl
21	42	56.8	696	3	US-09-907-942-91	Sequence 91, Appl
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37	42	56.8	696	3	US-09-907-925-91	Sequence 91, Appl
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54	42	56.8	696	3	US-09-903-925-91	Sequence 91, Appl
55	42	56.8	696	3	US-09-906-760A-91	Sequence 91, Appl
56	42	56.8	696	3	US-09-903-823-91	Sequence 91, Appl
57	42	56.8	696	3	US-09-907-652-91	Sequence 91, Appl
58	42	56.8	696	3	US-09-902-572A-91	Sequence 91, Appl
59	42	56.8	696	3	US-09-902-979-91	Sequence 91, Appl
60	42	56.8	696	3	US-09-905-125-91	Sequence 91, Appl
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62	42	56.8	696	3	US-09-905-449-91	Sequence 91, Appl
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66	42	56.8	696	3	US-09-906-777-91	Sequence 91, Appl
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69	42	56.8	696	3	US-09-904-766-91	Sequence 91, Appl
70	42	56.8	696	3	US-09-904-920A-91	Sequence 91, Appl
71	42	56.8	696	3	US-09-904-877A-91	Sequence 91, Appl
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73	42	56.8	696	3	US-09-906-618-91	Sequence 91, Appl
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75	42	56.8	696	3	US-09-904-805-91	Sequence 91, Appl
76	42	56.8	696	3	US-09-904-938A-91	Sequence 91, Appl
77	42	56.8	696	3	US-09-906-722A-91	Sequence 91, Appl
78	42	56.8	696	3	US-09-908-576-91	Sequence 91, Appl
79	42	56.8	696	4	US-10-028-072-354	Sequence 354, App
80	42	56.8	696	4	US-10-140-808-354	Sequence 354, App
81	42	56.8	696	4	US-10-121-049-354	Sequence 354, App
82	42	56.8	696	4	US-10-123-904-354	Sequence 354, App
83	42	56.8	696	4	US-10-140-470-354	Sequence 354, App
84	42	56.8	696	4	US-10-175-746-354	Sequence 354, App
85	42	56.8	696	4	US-10-176-918-354	Sequence 354, App
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91	42	56.8	696	4	US-10-142-419-354	Sequence 354, App
92	42	56.8	696	4	US-10-123-262-354	Sequence 354, App
93	42	56.8	696	4	US-10-142-423-354	Sequence 354, App
94	42	56.8	696	4	US-10-121-050-354	Sequence 354, App
95	42	56.8	696	4	US-10-141-755-354	Sequence 354, App
96	42	56.8	696	4	US-10-143-032-354	Sequence 354, App
97	42	56.8	696	4	US-10-123-108-354	Sequence 354, App
98	42	56.8	696	4	US-10-123-236-354	Sequence 354, App
99	42	56.8	696	4	US-10-123-261-354	Sequence 354, App
100	42	56.8	696	4	US-10-140-921-354	Sequence 354, App

ALIGNMENTS

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RESULT 1
US-10-769-514-17
; Sequence 17, Application US/10769514
; Publication No. US20040258695A1
; GENERAL INFORMATION:
;  APPLICANT: Schryvers, Anthony
;  TITLE OF INVENTION: Transferrin Binding Peptides and Uses Thereof
;  FILE REFERENCE: 028722-001
;  CURRENT APPLICATION NUMBER: US/10/769,514
;  CURRENT FILING DATE:  2004-01-30
;  PRIOR APPLICATION NUMBER: US 60/444,113
;  PRIOR FILING DATE: 2003-01-31
;  NUMBER OF SEQ ID NOS: 86
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 17
;   LENGTH: 14
;   TYPE: PRT
;   ORGANISM: M. catarrhalis
US-10-769-514-17

  Query Match             100.0%;  Score 74;  DB 5;  Length 14;
  Best Local Similarity   100.0%;  Pred. No. 2.9e-06;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||||
Db          1 MGYGMALSKINLHN 14


RESULT 2
US-10-769-514-15
; Sequence 15, Application US/10769514
; Publication No. US20040258695A1
; GENERAL INFORMATION:
;  APPLICANT: Schryvers, Anthony
;  TITLE OF INVENTION: Transferrin Binding Peptides and Uses Thereof
;  FILE REFERENCE: 028722-001
;  CURRENT APPLICATION NUMBER: US/10/769,514
;  CURRENT FILING DATE:  2004-01-30
;  PRIOR APPLICATION NUMBER: US 60/444,113
;  PRIOR FILING DATE: 2003-01-31
;  NUMBER OF SEQ ID NOS: 86
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 15
;   LENGTH: 15
;   TYPE: PRT
;   ORGANISM: M. catarrhalis
US-10-769-514-15

  Query Match             100.0%;  Score 74;  DB 5;  Length 15;
  Best Local Similarity   100.0%;  Pred. No. 3.2e-06;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||||
Db          1 MGYGMALSKINLHN 14


RESULT 3
US-10-769-514-16
; Sequence 16, Application US/10769514
; Publication No. US20040258695A1
; GENERAL INFORMATION:
;  APPLICANT: Schryvers, Anthony
;  TITLE OF INVENTION: Transferrin Binding Peptides and Uses Thereof
;  FILE REFERENCE: 028722-001
;  CURRENT APPLICATION NUMBER: US/10/769,514
;  CURRENT FILING DATE:  2004-01-30
;  PRIOR APPLICATION NUMBER: US 60/444,113
;  PRIOR FILING DATE: 2003-01-31
;  NUMBER OF SEQ ID NOS: 86
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 16
;   LENGTH: 15
;   TYPE: PRT
;   ORGANISM: M. catarrhalis
US-10-769-514-16

  Query Match             100.0%;  Score 74;  DB 5;  Length 15;
  Best Local Similarity   100.0%;  Pred. No. 3.2e-06;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||||
Db          2 MGYGMALSKINLHN 15


RESULT 4
US-10-425-115-367739
; Sequence 367739, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
;  APPLICANT: La Rosa, Thomas J.
;  APPLICANT:  Kovalic, David K.
;  APPLICANT:  Zhou, Yihua
;  APPLICANT:  Cao, Yongwei
;  TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
;  TITLE OF INVENTION:  Plants
;  FILE REFERENCE: 38-21(53222)B
;  CURRENT APPLICATION NUMBER: US/10/425,115
;  CURRENT FILING DATE:  2003-04-28
;  NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 367739
;   LENGTH: 64
;   TYPE: PRT
;   ORGANISM: Zea mays
;   FEATURE:
;   OTHER INFORMATION: Clone ID: MRT4577_98550C.1.pep
US-10-425-115-367739

  Query Match             59.5%;  Score 44;  DB 4;  Length 64;
  Best Local Similarity   72.7%;  Pred. No. 3.6;
  Matches    8;  Conservative    2;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINL 12
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Db         17 GYGYSISKINL 27


RESULT 5
US-10-424-599-250435
; Sequence 250435, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
;  APPLICANT: La Rosa Thomas J
;  APPLICANT:  Kovalic David K
;  APPLICANT:  Zhou Yihua
;  APPLICANT:  Cao Yongwei
;  TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
;  TITLE OF INVENTION:  Plants and Uses Thereof for Plant Improvement
;  FILE REFERENCE: 38-21(53223)B
;  CURRENT APPLICATION NUMBER: US/10/424,599
;  CURRENT FILING DATE:  2003-04-28
;  NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 250435
;   LENGTH: 71
;   TYPE: PRT
;   ORGANISM: Glycine max
;   FEATURE:
;   OTHER INFORMATION: Clone ID: PAT_MRT3847_68170C.1.pep
US-10-424-599-250435
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  Best Local Similarity   50.0%;  Pred. No. 6.1;
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Qy          1 MGYGMALSKINLHN 14
              :|||:   | |||:
Db         58 LGYGLTYDKENLHD 71


RESULT 6
US-10-424-233-19
; Sequence 19, Application US/10424233
; Publication No. US20030220263A1
; GENERAL INFORMATION:
;  APPLICANT: Bristol-Myers Squibb Company
;  TITLE OF INVENTION: NOVEL HUMAN LEUCINE-RICH REPEAT-CONTAINING PROTEINS SPECIFICALLY
;  TITLE OF INVENTION:  EXPRESSED IN THE NERVOUS SYSTEM
;  FILE REFERENCE: D0233 NP
;  CURRENT APPLICATION NUMBER: US/10/424,233
;  CURRENT FILING DATE:  2003-04-25
;  PRIOR APPLICATION NUMBER: U.S. 60/375,335
;  PRIOR FILING DATE: 2002-04-25
;  NUMBER OF SEQ ID NOS: 75
;  SOFTWARE: PatentIn version 3.2
; SEQ ID NO 19
;   LENGTH: 635
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-10-424-233-19

  Query Match             56.8%;  Score 42;  DB 4;  Length 635;
  Best Local Similarity   63.6%;  Pred. No. 98;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        432 GVSLSKLSLHN 442


RESULT 7
US-09-374-046A-132
; Sequence 132, Application US/09374046A
; Publication No. US20030096951A1
; GENERAL INFORMATION:
;  APPLICANT: Jacobs, Kenneth
;  APPLICANT: McCoy, John M.
;  APPLICANT: LaVallie, Edward R.
;  APPLICANT: Collins-Racie, Lisa A.
;  APPLICANT: Evans, Cheryl
;  APPLICANT: Merberg, David
;  APPLICANT: Treacy, Maurice
;  APPLICANT: Agostino, Michael J.
;  APPLICANT: Steininger II, Robert J.
;  APPLICANT: Spaulding, Vikki
;  APPLICANT: Wong, Gordon G.
;  APPLICANT: Clark, Hilary
;  APPLICANT: Fechtel, Kim
;  APPLICANT: Genetics Institute, Inc.
;  TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
;  FILE REFERENCE: GI 6075-83A
;  CURRENT APPLICATION NUMBER: US/09/374,046A
;  CURRENT FILING DATE: 1999-08-13
;  NUMBER OF SEQ ID NOS: 240
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 132
;   LENGTH: 695
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-374-046A-132

  Query Match             56.8%;  Score 42;  DB 3;  Length 695;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        492 GVSLSKLSLHN 502


RESULT 8
US-10-616-263-132
; Sequence 132, Application US/10616263
; Publication No. US20040038276A1
; GENERAL INFORMATION:
;  APPLICANT: Jacobs, Kenneth
;  APPLICANT:  McCoy, John M.
;  APPLICANT:  LaVallie, Edward R.
;  APPLICANT:  Collins-Racie, Lisa A.
;  APPLICANT:  Evans, Cheryl
;  APPLICANT:  Merberg, David
;  APPLICANT:  Treacy, Maurice
;  APPLICANT:  Agostino, Michael J.
;  APPLICANT:  Steininger II, Robert J.
;  APPLICANT:  Spaulding, Vikki
;  APPLICANT:  Wong, Gordon G.
;  APPLICANT:  Clark, Hilary
;  APPLICANT:  Fechtel, Kim
;  APPLICANT:  Genetics Institute, Inc.
;  TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
;  FILE REFERENCE: 00766.000103.5
;  CURRENT APPLICATION NUMBER: US/10/616,263
;  CURRENT FILING DATE:  2003-07-08
;  NUMBER OF SEQ ID NOS: 240
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 132
;   LENGTH: 695
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-10-616-263-132

  Query Match             56.8%;  Score 42;  DB 4;  Length 695;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        492 GVSLSKLSLHN 502


RESULT 9
US-09-909-320-91
; Sequence 91, Application US/09909320
; Patent No. US20020132240A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
```

```
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/909,320
;  CURRENT FILING DATE:  2002-01-04
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
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;  PRIOR FILING DATE: 1999-12-16
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;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-909-320-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
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Db        493 GVSLSKLSLHN 503


RESULT 10
US-09-909-088B-91
; Sequence 91, Application US/09909088B
; Patent No. US20020146709A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/909,088B
;  CURRENT FILING DATE:  2001-07-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-909-088B-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 11
US-09-905-291A-91
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; Sequence 91, Application US/09905291A
; Patent No. US20020160374A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,291A
; CURRENT FILING DATE: 2001-07-12
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-905-291A-91

Query Match 56.8%; Score 42; DB 3; Length 696;
Best Local Similarity 63.6%; Pred. No. 1.1e+02;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||
Db 493 GVSLSKLSLHN 503

RESULT 12
US-09-902-853-91
; Sequence 91, Application US/09902853
; Publication No. US20020192659A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/902,853
; CURRENT FILING DATE: 2001-07-10
; PRIOR APPLICATION NUMBER: US/09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16

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;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-902-853-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 13
US-09-907-824-91
; Sequence 91, Application US/09907824
; Publication No. US20020197671A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/907,824
;  CURRENT FILING DATE:  2001-07-17
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-907-824-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
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Db        493 GVSLSKLSLHN 503


RESULT 14
US-09-907-841-91
; Sequence 91, Application US/09907841
; Publication No. US20020198366A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/907,841
;  CURRENT FILING DATE:  2001-11-20
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
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;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  Remaining Prior Application data removed - See File Wrapper or PALM.
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;    LENGTH: 696
;    TYPE: PRT
;    ORGANISM: Homo sapiens
US-09-907-841-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 15
US-09-904-011-91
; Sequence 91, Application US/09904011
; Publication No. US20030003530A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,011
;  CURRENT FILING DATE:  2001-07-11
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;    LENGTH: 696
;    TYPE: PRT
;    ORGANISM: Homo Sapien
US-09-904-011-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 16
US-09-903-640-91
; Sequence 91, Application US/09903640
; Publication No. US20030017463A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
```

```
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/903,640
;  CURRENT FILING DATE:  2001-07-11
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-903-640-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 17
US-09-908-093-91
; Sequence 91, Application US/09908093
; Publication No. US20030017498A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/908,093
;  CURRENT FILING DATE:  2001-07-17
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-908-093-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 18
US-09-906-742-91
; Sequence 91, Application US/09906742
; Publication No. US20030023054A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
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;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,742
;  CURRENT FILING DATE:  2001-07-16
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-906-742-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 19
US-09-906-838-91
; Sequence 91, Application US/09906838
; Publication No. US20030027143A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,838
;  CURRENT FILING DATE:  2001-07-16
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-906-838-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
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Db        493 GVSLSKLSLHN 503


RESULT 20
US-09-907-613-91
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; Sequence 91, Application US/09907613
; Publication No. US20030027145A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,613
; CURRENT FILING DATE: 2001-07-17
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-907-613-91

Query Match 56.8%; Score 42; DB 3; Length 696;
Best Local Similarity 63.6%; Pred. No. 1.1e+02;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

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Qy          4 GMALSKINLHN 14
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Db        493 GVSLSKLSLHN 503
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RESULT 21
US-09-907-942-91
; Sequence 91, Application US/09907942
; Publication No. US20030027146A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,942
; CURRENT FILING DATE: 2002-01-22
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16

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;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-907-942-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 22
US-09-904-859-91
; Sequence 91, Application US/09904859
; Publication No. US20030036060A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,859
;  CURRENT FILING DATE:  2001-07-12
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
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;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-904-859-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 23
US-09-909-204-91
; Sequence 91, Application US/09909204
; Publication No. US20030036061A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/909,204
;  CURRENT FILING DATE:  2001-07-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
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;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
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;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-909-204-91

  Query Match            56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 24
US-09-904-820-91
; Sequence 91, Application US/09904820
; Publication No. US20030036094A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,820
;  CURRENT FILING DATE:  2001-07-13
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-904-820-91

  Query Match            56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 25
US-09-904-786-91
; Sequence 91, Application US/09904786
; Publication No. US20030039969A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
```

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;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,786
;  CURRENT FILING DATE:  2001-07-12
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-904-786-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 26
US-09-906-646-91
; Sequence 91, Application US/09906646
; Publication No. US20030039971A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,646
;  CURRENT FILING DATE:  2002-01-22
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-906-646-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 27
US-09-906-700-91
; Sequence 91, Application US/09906700
; Publication No. US20030039972A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
```

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;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/906,700
;  CURRENT FILING DATE:  2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-906-700-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 28
US-09-903-786-91
; Sequence 91, Application US/09903786
; Publication No. US20030044793A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/903,786
;  CURRENT FILING DATE:  2001-07-11
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-903-786-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
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  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 29
US-09-902-903-91
; Sequence 91, Application US/09902903
; Publication No. US20030044839A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/902,903
;  CURRENT FILING DATE:  2001-07-10
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-902-903-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 30
US-09-903-749A-91
; Sequence 91, Application US/09903749A
; Publication No. US20030045693A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/903,749A
;  CURRENT FILING DATE:  2001-07-11
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
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;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-903-749A-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 31
US-09-904-119-91
; Sequence 91, Application US/09904119
; Publication No. US20030049621A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,119
;  CURRENT FILING DATE:  2001-07-11
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-904-119-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 32
US-09-904-956-91
; Sequence 91, Application US/09904956
; Publication No. US20030049622A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
```

```
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,956
;  CURRENT FILING DATE:  2001-07-12
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-904-956-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 33
US-09-902-736-91
; Sequence 91, Application US/09902736
; Publication No. US20030049676A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/902,736
;  CURRENT FILING DATE:  2001-07-10
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-902-736-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 34
US-09-907-794-91
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; Sequence 91, Application US/09907794
; Publication No. US20030049677A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/907,794
; CURRENT FILING DATE: 2001-07-17
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/28565
; PRIOR FILING DATE: 1999-12-02
; PRIOR APPLICATION NUMBER: PCT/US99/30095
; PRIOR FILING DATE: 1999-12-16
; PRIOR APPLICATION NUMBER: PCT/US99/30911
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US99/30999
; PRIOR FILING DATE: 1999-12-20
; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-907-794-91

Query Match 56.8%; Score 42; DB 3; Length 696;
Best Local Similarity 63.6%; Pred. No. 1.1e+02;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 4 GMALSKINLHN 14
|::|||::|||
Db 493 GVSLSKLSLHN 503

RESULT 35
US-09-903-943-91
; Sequence 91, Application US/09903943
; Publication No. US20030054349A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/903,943
; CURRENT FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
; PRIOR FILING DATE: 1999-07-28
; PRIOR APPLICATION NUMBER: PCT/US99/20594
; PRIOR FILING DATE: 1999-09-08
; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/21547
; PRIOR FILING DATE: 1999-09-15
; PRIOR APPLICATION NUMBER: PCT/US99/23089
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: PCT/US99/28214
; PRIOR FILING DATE: 1999-11-29
; PRIOR APPLICATION NUMBER: PCT/US99/28313
; PRIOR FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: PCT/US99/28564
; PRIOR FILING DATE: 1999-12-02

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;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-903-943-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 36
US-09-904-462-91
; Sequence 91, Application US/09904462
; Publication No. US20030054351A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/904,462
;  CURRENT FILING DATE:  2001-07-13
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-904-462-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 37
US-09-907-925-91
; Sequence 91, Application US/09907925
; Publication No. US20030054352A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
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;  CURRENT APPLICATION NUMBER: US/09/907,925
;  CURRENT FILING DATE:  2001-07-17
;  PRIOR APPLICATION NUMBER: 09/665,350
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-09-907-925-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
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Db        493 GVSLSKLSLHN 503


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; Sequence 91, Application US/09902692
; Publication No. US20030054400A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
;  APPLICANT:  Botstein, David
;  APPLICANT:  Desnoyers, Luc
;  APPLICANT:  Eaton, Dan L.
;  APPLICANT:  Ferrara, Napoleone
;  APPLICANT:  Filvaroff, Ellen
;  APPLICANT:  Fong, Sherman
;  APPLICANT:  Gao, Wei-Qiang
;  APPLICANT:  Gerber, Hanspeter
;  APPLICANT:  Gerritsen, Mary E.
;  APPLICANT:  Goddard, A.
;  APPLICANT:  Godowski, Paul J.
;  APPLICANT:  Grimaldi, Christopher J.
;  APPLICANT:  Gurney, Austin L.
;  APPLICANT:  Hillan, Kenneth, J.
;  APPLICANT:  Kljavin, Ivar J.
;  APPLICANT:  Mather, Jennie P.
;  APPLICANT:  Pan, James
;  APPLICANT:  Paoni, Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William, I.
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: 10466-14
;  CURRENT APPLICATION NUMBER: US/09/902,692
;  CURRENT FILING DATE:  2001-07-10
;  PRIOR APPLICATION NUMBER: PCT/US00/04414
;  PRIOR FILING DATE: 2000-02-22
;  PRIOR APPLICATION NUMBER: US 60/143,048
;  PRIOR FILING DATE: 1999-07-07
;  PRIOR APPLICATION NUMBER: US 60/145,698
;  PRIOR FILING DATE: 1999-07-26
;  PRIOR APPLICATION NUMBER: US 60/146,222
;  PRIOR FILING DATE: 1999-07-28
;  PRIOR APPLICATION NUMBER: PCT/US99/20594
;  PRIOR FILING DATE: 1999-09-08
;  PRIOR APPLICATION NUMBER: PCT/US99/20944
;  PRIOR FILING DATE: 1999-09-13
;  PRIOR APPLICATION NUMBER: PCT/US99/21090
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/21547
;  PRIOR FILING DATE: 1999-09-15
;  PRIOR APPLICATION NUMBER: PCT/US99/23089
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: PCT/US99/28214
;  PRIOR FILING DATE: 1999-11-29
;  PRIOR APPLICATION NUMBER: PCT/US99/28313
;  PRIOR FILING DATE: 1999-11-30
;  PRIOR APPLICATION NUMBER: PCT/US99/28564
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/28565
;  PRIOR FILING DATE: 1999-12-02
;  PRIOR APPLICATION NUMBER: PCT/US99/30095
;  PRIOR FILING DATE: 1999-12-16
;  PRIOR APPLICATION NUMBER: PCT/US99/30911
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US99/30999
;  PRIOR FILING DATE: 1999-12-20
;  PRIOR APPLICATION NUMBER: PCT/US00/00219
;  PRIOR FILING DATE: 2000-01-05
;  NUMBER OF SEQ ID NOS: 423
; SEQ ID NO 91
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-902-692-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

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Db        493 GVSLSKLSLHN 503


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; Sequence 91, Application US/09903520
; Publication No. US20030054401A1
; GENERAL INFORMATION:
;  APPLICANT: Genentech, Inc.
;  APPLICANT:  Ashkenazi, Avi
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; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/903,520
; CURRENT FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: 09/665,350
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
; PRIOR APPLICATION NUMBER: US 60/143,048
; PRIOR FILING DATE: 1999-07-07
; PRIOR APPLICATION NUMBER: US 60/145,698
; PRIOR FILING DATE: 1999-07-26
; PRIOR APPLICATION NUMBER: US 60/146,222
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; PRIOR APPLICATION NUMBER: PCT/US99/20944
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: PCT/US99/21090
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; PRIOR FILING DATE: 1999-11-29
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; PRIOR APPLICATION NUMBER: PCT/US00/00219
; PRIOR FILING DATE: 2000-01-05
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; ORGANISM: Homo Sapien
US-09-903-520-91

Query Match 56.8%; Score 42; DB 3; Length 696;
Best Local Similarity 63.6%; Pred. No. 1.1e+02;

Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

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; Sequence 91, Application US/09905056
; Publication No. US20030054441A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi
; APPLICANT: Botstein, David
; APPLICANT: Desnoyers, Luc
; APPLICANT: Eaton, Dan L.
; APPLICANT: Ferrara, Napoleone
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Gao, Wei-Qiang
; APPLICANT: Gerber, Hanspeter
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, A.
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Hillan, Kenneth, J.
; APPLICANT: Kljavin, Ivar J.
; APPLICANT: Mather, Jennie P.
; APPLICANT: Pan, James
; APPLICANT: Paoni, Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William, I.
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: 10466-14
; CURRENT APPLICATION NUMBER: US/09/905,056
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; PRIOR APPLICATION NUMBER: PCT/US00/04414
; PRIOR FILING DATE: 2000-02-22
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; PRIOR APPLICATION NUMBER: PCT/US99/30095
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;  PRIOR APPLICATION NUMBER: PCT/US00/00219
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; SEQ ID NO 91
;   LENGTH: 696
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;   ORGANISM: Homo sapiens
US-09-905-056-91

  Query Match             56.8%;  Score 42;  DB 3;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.1e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

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GenCore version 5.1.8

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Searched: 250354 seqs, 46948837 residues

Total number of hits satisfying chosen parameters: 250354

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Post-processing: Minimum Match 0%
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Listing first 100 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2	42	56.8	696	9	US-10-511-538-231	Sequence 231, App
3	42	56.8	696	9	US-10-973-115B-354	Sequence 354, App
4	42	56.8	696	9	US-10-137-873A-354	Sequence 354, App
5	42	56.8	696	9	US-10-152-370-354	Sequence 354, App
6	42	56.8	696	11	US-11-290-153-354	Sequence 354, App
7	37	50.0	310	11	US-11-079-463-9199	Sequence 9199, Ap
8	36.5	49.3	350	11	US-11-108-528-54	Sequence 54, Appl
9	36.5	49.3	351	11	US-11-108-528-52	Sequence 52, Appl
10	36	48.6	499	11	US-11-225-903-17	Sequence 17, Appl
11	36	48.6	532	11	US-11-045-004-2431	Sequence 2431, Ap
12	35	47.3	175	11	US-11-172-740-1344	Sequence 1344, Ap
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16	35	47.3	341	11	US-11-188-298-11417	Sequence 11417, A
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26	35	47.3	1588	9	US-10-995-561-527	Sequence 527, App
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33	34	45.9	445	11	US-11-188-298-7996	Sequence 7996, Ap
34	34	45.9	467	11	US-11-079-463-7672	Sequence 7672, Ap
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37	34	45.9	662	9	US-10-915-002-308	Sequence 308, App
38	34	45.9	724	11	US-11-072-512-2224	Sequence 2224, Ap
39	34	45.9	845	11	US-11-147-047-46	Sequence 46, Appl
40	34	45.9	845	11	US-11-264-096-483	Sequence 483, App
41	33.5	45.3	533	11	US-11-188-298-20107	Sequence 20107, A
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43	33	44.6	174	11	US-11-096-568A-739	Sequence 739, App
44	33	44.6	198	11	US-11-188-298-2235	Sequence 2235, Ap
45	33	44.6	232	11	US-11-096-568A-26655	Sequence 26655, A
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47	33	44.6	283	9	US-10-784-004-386	Sequence 386, App
48	33	44.6	283	9	US-10-784-004-707	Sequence 707, App
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55	33	44.6	338	9	US-10-520-820-6	Sequence 6, Appli
56	33	44.6	356	11	US-11-096-568A-26653	Sequence 26653, A
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58	33	44.6	363	11	US-11-188-298-22281	Sequence 22281, A
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60	33	44.6	376	11	US-11-098-686-11320	Sequence 11320, A
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63	33	44.6	379	11	US-11-096-568A-15418	Sequence 15418, A
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69	33	44.6	479	11	US-11-096-568A-28911	Sequence 28911, A
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77	33	44.6	575	11	US-11-188-298-14037	Sequence 14037, A
78	33	44.6	583	11	US-11-087-099-9797	Sequence 9797, Ap
79	33	44.6	610	9	US-10-793-626-482	Sequence 482, App
80	33	44.6	614	11	US-11-079-463-10331	Sequence 10331, A
81	33	44.6	625	11	US-11-188-298-19611	Sequence 19611, A
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83	33	44.6	642	11	US-11-072-175-172	Sequence 172, App
84	33	44.6	665	11	US-11-045-004-909	Sequence 909, App
85	33	44.6	731	11	US-11-188-298-9427	Sequence 9427, Ap
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87	33	44.6	732	11	US-11-188-298-668	Sequence 668, App
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92	33	44.6	773	11	US-11-052-554A-341	Sequence 341, App
93	33	44.6	845	11	US-11-188-298-14852	Sequence 14852, A
94	33	44.6	855	11	US-11-087-099-12274	Sequence 12274, A

95	33	44.6	918	11	US-11-024-959-492	Sequence 492, App
96	33	44.6	925	11	US-11-079-463-6249	Sequence 6249, Ap
97	33	44.6	1445	11	US-11-124-367A-478	Sequence 478, App
98	33	44.6	1464	9	US-10-912-971-4	Sequence 4, Appli
99	33	44.6	1464	11	US-11-076-074-1	Sequence 1, Appli
100	33	44.6	1464	11	US-11-124-367A-262	Sequence 262, App

ALIGNMENTS

RESULT 1
US-10-131-826A-354
; Sequence 354, Application US/10131826A
; Publication No. US20050245730A1
; GENERAL INFORMATION:
; APPLICANT: Baker,Kevin P.
; APPLICANT: Beresini,Maureen
; APPLICANT: DeForge,Laura
; APPLICANT: Desnoyers,Luc
; APPLICANT: Filvaroff,Ellen
; APPLICANT: Gao,Wei-Qiang
; APPLICANT: Gerritsen,Mary E.
; APPLICANT: Goddard,Audrey
; APPLICANT: Godowski,Paul J.
; APPLICANT: Gurney,Austin L.
; APPLICANT: Sherwood,Steven
; APPLICANT: Smith,Victoria
; APPLICANT: Stewart,Timothy A.
; APPLICANT: Tumas,Daniel
; APPLICANT: Watanabe,Colin K
; APPLICANT: Wood,William
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; TITLE OF INVENTION: ACIDS ENCODING THE SAME
; FILE REFERENCE: P3330R1C128
; CURRENT APPLICATION NUMBER: US/10/131,826A
; CURRENT FILING DATE: 2002-04-24
; PRIOR APPLICATION NUMBER: 60/049911
; PRIOR FILING DATE: 1997-06-18
; PRIOR APPLICATION NUMBER: 60/056974
; PRIOR FILING DATE: 1997-08-26
; PRIOR APPLICATION NUMBER: 60/059113
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059115
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059117
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059122
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059184
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059263
; PRIOR FILING DATE: 1997-09-18
; PRIOR APPLICATION NUMBER: 60/059352
; PRIOR FILING DATE: 1997-09-19
; PRIOR APPLICATION NUMBER: 60/059588
; PRIOR FILING DATE: 1997-09-19
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 354
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-131-826A-354

Query Match 56.8%; Score 42; DB 9; Length 696;
Best Local Similarity 63.6%; Pred. No. 10;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
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RESULT 2
US-10-511-538-231
; Sequence 231, Application US/10511538
; Publication No. US20060026700A1
; GENERAL INFORMATION:
; APPLICANT: OriGene Technologies, Inc
; TITLE OF INVENTION: TISSUE SPECIFIC GENES AND GENE CLUSTERS
; FILE REFERENCE: 16U 200 PCT
; CURRENT APPLICATION NUMBER: US/10/511,538
; CURRENT FILING DATE: 2004-10-18
; PRIOR APPLICATION NUMBER: US 60/372,669
; PRIOR FILING DATE: 2002-04-16
; PRIOR APPLICATION NUMBER: US 60/411,882
; PRIOR FILING DATE: 2002-09-20
; PRIOR APPLICATION NUMBER: US 60/424,336
; PRIOR FILING DATE: 2002-11-07
; PRIOR APPLICATION NUMBER: US 60/374,823
; PRIOR FILING DATE: 2002-04-24
; PRIOR APPLICATION NUMBER: US 60/376,558
; PRIOR FILING DATE: 2002-05-01
; PRIOR APPLICATION NUMBER: US 60/381,366
; PRIOR FILING DATE: 2002-05-20
; PRIOR APPLICATION NUMBER: US 60/403,648
; PRIOR FILING DATE: 2002-08-16
; NUMBER OF SEQ ID NOS: 344
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 231
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-511-538-231

Query Match 56.8%; Score 42; DB 9; Length 696;
Best Local Similarity 63.6%; Pred. No. 10;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
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RESULT 3
US-10-973-115B-354
; Sequence 354, Application US/10973115B
; Publication No. US20060040351A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
; APPLICANT: Beresini, Maureen
; APPLICANT: DeForge, Laura
; APPLICANT: Desnoyers, Luc
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gao, Wei-Quiang
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul
; APPLICANT: Gurney, Austin L.
; APPLICANT: Sherwood, Steven
; APPLICANT: Smith, Victoria
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ACIDS ENCODIN
; TITLE OF INVENTION: SAME
; FILE REFERENCE: 39870-3330R1C300C1
; CURRENT APPLICATION NUMBER: US/10/973,115B
; CURRENT FILING DATE: 2004-10-22
; PRIOR APPLICATION NUMBER: US 10/145,747
; PRIOR FILING DATE: 2002-05-14
; PRIOR APPLICATION NUMBER: US 10/028,072

; PRIOR FILING DATE: 2001-12-19
; PRIOR APPLICATION NUMBER: PCT/US00/32678
; PRIOR FILING DATE: 2000-12-01
; PRIOR APPLICATION NUMBER: US 09/581,742
; PRIOR FILING DATE: 2000-06-16
; PRIOR APPLICATION NUMBER: PCT/US00/05746
; PRIOR FILING DATE: 2000-03-02
; PRIOR APPLICATION NUMBER: US 60/135,736
; PRIOR FILING DATE: 1999-05-25
; PRIOR APPLICATION NUMBER: US 60/123,090
; PRIOR FILING DATE: 1999-03-05
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 354
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-973-115B-354

Query Match 56.8%; Score 42; DB 9; Length 696;
Best Local Similarity 63.6%; Pred. No. 10;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
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RESULT 4
US-10-137-873A-354
; Sequence 354, Application US/10137873A
; Publication No. US20060084138A1
; GENERAL INFORMATION:
; APPLICANT: Baker,Kevin P.
; APPLICANT: Beresini,Maureen
; APPLICANT: DeForge,Laura
; APPLICANT: Desnoyers,Luc
; APPLICANT: Filvaroff,Ellen
; APPLICANT: Gao,Wei-Qiang
; APPLICANT: Gerritsen,Mary E.
; APPLICANT: Goddard,Audrey
; APPLICANT: Godowski,Paul J.
; APPLICANT: Gurney,Austin L.
; APPLICANT: Sherwood,Steven
; APPLICANT: Smith,Victoria
; APPLICANT: Stewart,Timothy A.
; APPLICANT: Tumas,Daniel
; APPLICANT: Watanabe,Colin K
; APPLICANT: Wood,William
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; TITLE OF INVENTION: ACIDS ENCODING THE SAME
; FILE REFERENCE: P3330R1C149
; CURRENT APPLICATION NUMBER: US/10/137,873A
; CURRENT FILING DATE: 2002-04-23
; PRIOR APPLICATION NUMBER: 60/049911
; PRIOR FILING DATE: 1997-06-18
; PRIOR APPLICATION NUMBER: 60/056974
; PRIOR FILING DATE: 1997-08-26
; PRIOR APPLICATION NUMBER: 60/059113
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059115
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059117
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059122
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059184
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059263
; PRIOR FILING DATE: 1997-09-18
; PRIOR APPLICATION NUMBER: 60/059352
; PRIOR FILING DATE: 1997-09-19
; PRIOR APPLICATION NUMBER: 60/059588
; PRIOR FILING DATE: 1997-09-19
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 354
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-137-873A-354

Query Match 56.8%; Score 42; DB 9; Length 696;
Best Local Similarity 63.6%; Pred. No. 10;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 5
US-10-152-370-354
; Sequence 354, Application US/10152370
; Publication No. US20060084139A1
; GENERAL INFORMATION:
; APPLICANT: Baker,Kevin P.
; APPLICANT: Beresini,Maureen
; APPLICANT: DeForge,Laura
; APPLICANT: Desnoyers,Luc
; APPLICANT: Filvaroff,Ellen
; APPLICANT: Gao,Wei-Qiang
; APPLICANT: Gerritsen,Mary E.
; APPLICANT: Goddard,Audrey
; APPLICANT: Godowski,Paul J.
; APPLICANT: Gurney,Austin L.
; APPLICANT: Sherwood,Steven
; APPLICANT: Smith,Victoria
; APPLICANT: Stewart,Timothy A.
; APPLICANT: Tumas,Daniel
; APPLICANT: Watanabe,Colin K
; APPLICANT: Wood,William
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; TITLE OF INVENTION: ACIDS ENCODING THE SAME
; FILE REFERENCE: P3330R1C407
; CURRENT APPLICATION NUMBER: US/10/152,370
; CURRENT FILING DATE: 2002-05-21
; Prior Application removed - See file Wrapper or Palm
; NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 354
; LENGTH: 696
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-152-370-354

Query Match 56.8%; Score 42; DB 9; Length 696;
Best Local Similarity 63.6%; Pred. No. 10;
Matches 7; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

```
Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
```

RESULT 6
US-11-290-153-354
; Sequence 354, Application US/11290153
; Publication No. US20060073568A1
; GENERAL INFORMATION:
; APPLICANT: Baker,Kevin P.
; APPLICANT: Beresini,Maureen
; APPLICANT: DeForge,Laura
; APPLICANT: Desnoyers,Luc
; APPLICANT: Filvaroff,Ellen
; APPLICANT: Gao,Wei-Qiang

```
;  APPLICANT:  Gerritsen,Mary E.
;  APPLICANT:  Goddard,Audrey
;  APPLICANT:  Godowski,Paul J.
;  APPLICANT:  Gurney,Austin L.
;  APPLICANT:  Sherwood,Steven
;  APPLICANT:  Smith,Victoria
;  APPLICANT:  Stewart,Timothy A.
;  APPLICANT:  Tumas,Daniel
;  APPLICANT:  Watanabe,Colin K
;  APPLICANT:  Wood,William
;  APPLICANT:  Zhang, Zemin
;  TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
;  TITLE OF INVENTION:  ACIDS ENCODING THE SAME
;  FILE REFERENCE: P3330R1C321
;  CURRENT APPLICATION NUMBER: US/11/290,153
;  CURRENT FILING DATE:  2005-11-30
;  PRIOR APPLICATION NUMBER: US/10/146,728
;  PRIOR FILING DATE: 2002-05-15
;  PRIOR APPLICATION NUMBER: 60/049911
;  PRIOR FILING DATE: 1997-06-18
;  PRIOR APPLICATION NUMBER: 60/056974
;  PRIOR FILING DATE: 1997-08-26
;  PRIOR APPLICATION NUMBER: 60/059113
;  PRIOR FILING DATE: 1997-09-17
;  PRIOR APPLICATION NUMBER: 60/059115
;  PRIOR FILING DATE: 1997-09-17
;  PRIOR APPLICATION NUMBER: 60/059117
;  PRIOR FILING DATE: 1997-09-17
;  PRIOR APPLICATION NUMBER: 60/059122
;  PRIOR FILING DATE: 1997-09-17
;  PRIOR APPLICATION NUMBER: 60/059184
;  PRIOR FILING DATE: 1997-09-17
;  PRIOR APPLICATION NUMBER: 60/059263
;  PRIOR FILING DATE: 1997-09-18
;  PRIOR APPLICATION NUMBER: 60/059352
;  PRIOR FILING DATE: 1997-09-19
;  Remaining Prior Application data removed - See File Wrapper or PALM.
;  NUMBER OF SEQ ID NOS: 550
; SEQ ID NO 354
;   LENGTH: 696
;   TYPE: PRT
;   ORGANISM: Homo Sapien
US-11-290-153-354

  Query Match             56.8%;  Score 42;  DB 11;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 10;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 7
US-11-079-463-9199
; Sequence 9199, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
;  APPLICANT: Gary L. Breton
;  TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES FRA
;  TITLE OF INVENTION:  FOR DIAGNOSTICS AND THERAPEUTICS
;  FILE REFERENCE: PATH00-03DIV2
;  CURRENT APPLICATION NUMBER: US/11/079,463
;  CURRENT FILING DATE:  2005-03-14
;  PRIOR APPLICATION NUMBER: US 60/128,705
;  PRIOR FILING DATE: 1999-04-09
;  PRIOR APPLICATION NUMBER: US 09/540,209
;  PRIOR FILING DATE: 2000-04-04
;  NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 9199
;   LENGTH: 310
;   TYPE: PRT
;   ORGANISM: B.fragilis
US-11-079-463-9199

  Query Match             50.0%;  Score 37;  DB 11;  Length 310;
  Best Local Similarity   54.5%;  Pred. No. 35;
  Matches    6;  Conservative    3;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          3 YGMALSKINLH 13
              ||:||  ::||
Db         65 YGLALEVVDLH 75


RESULT 8
US-11-108-528-54
; Sequence 54, Application US/11108528
; Publication No. US20050261189A1
; GENERAL INFORMATION:
;  APPLICANT: Larsen, Glenn
;  APPLICANT:  Marvin, Martha
;  APPLICANT:  Li, Dean Y.
;  APPLICANT:  Wang, Elizabeth
;  APPLICANT:  Chen, C. M. Amy
;  APPLICANT:  Shamah, Steven M.
;  TITLE OF INVENTION: METHODS OF PROMOTING CARDIAC CELL
;  TITLE OF INVENTION:  PROLIFERATION
;  FILE REFERENCE: HYDR-P01-041
;  CURRENT APPLICATION NUMBER: US/11/108,528
;  CURRENT FILING DATE:  2005-04-18
;  PRIOR APPLICATION NUMBER: US 60/563,137
;  PRIOR FILING DATE: 2004-04-16
;  PRIOR APPLICATION NUMBER: US 60/598,368
;  PRIOR FILING DATE: 2004-08-02
;  NUMBER OF SEQ ID NOS: 86
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 54
;   LENGTH: 350
;   TYPE: PRT
;   ORGANISM: Mouse
US-11-108-528-54

  Query Match             49.3%;  Score 36.5;  DB 11;  Length 350;
  Best Local Similarity   31.0%;  Pred. No. 49;
  Matches    9;  Conservative    4;  Mismatches    1;  Indels   15;  Gaps    1;

Qy          1 MGYGMALSK---------------INLHN 14
              :|:| |:||               :||||
Db        135 VGFGEAISKQFVDALETGQDARAAMNLHN 163


RESULT 9
US-11-108-528-52
; Sequence 52, Application US/11108528
; Publication No. US20050261189A1
; GENERAL INFORMATION:
;  APPLICANT: Larsen, Glenn
;  APPLICANT:  Marvin, Martha
;  APPLICANT:  Li, Dean Y.
;  APPLICANT:  Wang, Elizabeth
;  APPLICANT:  Chen, C. M. Amy
;  APPLICANT:  Shamah, Steven M.
;  TITLE OF INVENTION: METHODS OF PROMOTING CARDIAC CELL
;  TITLE OF INVENTION:  PROLIFERATION
;  FILE REFERENCE: HYDR-P01-041
;  CURRENT APPLICATION NUMBER: US/11/108,528
;  CURRENT FILING DATE:  2005-04-18
;  PRIOR APPLICATION NUMBER: US 60/563,137
;  PRIOR FILING DATE: 2004-04-16
;  PRIOR APPLICATION NUMBER: US 60/598,368
;  PRIOR FILING DATE: 2004-08-02
;  NUMBER OF SEQ ID NOS: 86
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 52
;   LENGTH: 351
;   TYPE: PRT
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;   ORGANISM: Homo sapiens
US-11-108-528-52

  Query Match             49.3%;  Score 36.5;  DB 11;  Length 351;
  Best Local Similarity   31.0%;  Pred. No. 49;
  Matches    9;  Conservative    4;  Mismatches    1;  Indels   15;  Gaps    1;

Qy          1 MGYGMALSK---------------INLHN 14
              :|:| |:||               :||||
Db        136 VGPGEAISKQFVDALETGQDARAAMNLHN 164


RESULT 10
US-11-225-903-17
; Sequence 17, Application US/11225903
; Publication No. US20060059582A1
; GENERAL INFORMATION:
;  APPLICANT: Jankowski, Boris
;  APPLICANT:  Feldmann, Kenneth A.
;  APPLICANT:  Bobzin, Steven Craig
;  TITLE OF INVENTION: Modulation of Amino Acid and Sugar Content in Plants
;  FILE REFERENCE: 11696-137001/WO1
;  CURRENT APPLICATION NUMBER: US/11/225,903
;  CURRENT FILING DATE:  2005-09-13
;  PRIOR APPLICATION NUMBER: 60/610,356
;  PRIOR FILING DATE: 2004-9-14
;  NUMBER OF SEQ ID NOS: 28
; SEQ ID NO 17
;   LENGTH: 499
;   TYPE: PRT
;   ORGANISM: Arabidopsis thaliana
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)..(499)
;   OTHER INFORMATION: Public GI no. 11994438
US-11-225-903-17

  Query Match             48.6%;  Score 36;  DB 11;  Length 499;
  Best Local Similarity   60.0%;  Pred. No. 92;
  Matches    6;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          3 YGMALSKINL 12
              |||| :: ||:
Db        468 YGMAIADINM 477


RESULT 11
US-11-045-004-2431
; Sequence 2431, Application US/11045004
; Publication No. US20060078901A1
; GENERAL INFORMATION:
;  APPLICANT: BUCHRIESER, CARMEN
;  APPLICANT:  FRANGEUL, LIONEL
;  APPLICANT:  COUVE, ELISABETH
;  APPLICANT:  RUSNIOK, CHRISTOPHE
;  APPLICANT:  FSIHI, HAFIDA
;  APPLICANT:  DEHOUX, PIERRE
;  APPLICANT:  DUSSURGET, OLIVIER
;  APPLICANT:  CHETOUANI, FARID
;  APPLICANT:  NEDJARI, HAFED
;  APPLICANT:  GLASER, PHILIPPE
;  APPLICANT:  KUNST, FRANCK
;  APPLICANT:  COSSART, PASCALE
;  APPLICANT:  DANIELS, JUSTIN
;  APPLICANT:  GOEBEI, WERNER
;  APPLICANT:  KREFT, JURGEN
;  APPLICANT:  KUHN, MICHAEL
;  APPLICANT:  NG, EVA
;  APPLICANT:  VAZQUEZ-BOLAND, ANTONIO
;  APPLICANT:  DOMINGUEZ-BERNAL, GUSTAVO
;  APPLICANT:  GARRIDO-GARCIA, PATRICIA
;  APPLICANT:  TIERREZ-MARTINEZ, ALBERTO
;  APPLICANT:  AMEND, ALEXANDRA
;  APPLICANT:  CHAKRABORTY, TRINAD
;  APPLICANT:  DOMANN, EUGEN
;  APPLICANT:  HAIN, THORSTEN
;  APPLICANT:  BERCHE, PATRICK
;  APPLICANT:  CHARBIT, ALAIN
;  APPLICANT:  DURANT, LIONEL
;  APPLICANT:  PEREZ-DIAZ, JOSE-CLAUDIO
;  APPLICANT:  BAQUERO, FERNANDO
;  APPLICANT:  GARCIA DEL PORTILLO, FRANCISCO
;  APPLICANT:  GOMEZ-LOPEZ, NURIA
;  APPLICANT:  MADUENIO, ENCARNA
;  APPLICANT:  PABLOS, BETRIZ DE
;  APPLICANT:  WEHLAND, JURGEN
;  APPLICANT:  KARST, UWE
;  APPLICANT:  ENTIAN, KARL-DIETER
;  APPLICANT:  HAUF, JORG
;  APPLICANT:  ROSE, MATTHIAS
;  APPLICANT:  VOSS, HAMUT
;  TITLE OF INVENTION: LISTERIA MONOCYTOGENES GENOME, POLYPEPTIDES AND USES
;  FILE REFERENCE: 05394.0018-02
;  CURRENT APPLICATION NUMBER: US/11/045,004
;  CURRENT FILING DATE:  2005-01-28
;  PRIOR APPLICATION NUMBER: 10/637,657
;  PRIOR FILING DATE: 2003-08-11
;  PRIOR APPLICATION NUMBER: 10/257,023
;  PRIOR FILING DATE: 2002-10-08
;  PRIOR APPLICATION NUMBER: PCT/FR01/01118
;  PRIOR FILING DATE: 2001-04-11
;  PRIOR APPLICATION NUMBER: FR 00/04,629
;  PRIOR FILING DATE: 2000-04-11
;  NUMBER OF SEQ ID NOS: 2854
;  SOFTWARE: PatentIn version 3.3
; SEQ ID NO 2431
;   LENGTH: 532
;   TYPE: PRT
;   ORGANISM: Listeria monocytogenes
US-11-045-004-2431

  Query Match             48.6%;  Score 36;  DB 11;  Length 532;
  Best Local Similarity   54.5%;  Pred. No. 99;
  Matches    6;  Conservative    3;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          3 YGMALSKINLH 13
              || || ::||:
Db         71 YGQALERLNLN 81


RESULT 12
US-11-172-740-1344
; Sequence 1344, Application US/11172740
; Publication No. US20060057724A1
; GENERAL INFORMATION:
;  APPLICANT: MASCIA, Peter
;  APPLICANT:  ALEXANDROV, Nickolai
;  APPLICANT:  BROVER, Vyacheslav
;  TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
;  TITLE OF INVENTION:  PLANT CHARACTERISTICS AND PHENOTYPES
;  FILE REFERENCE: 2750-1602PUS2
;  CURRENT APPLICATION NUMBER: US/11/172,740
;  CURRENT FILING DATE:  2005-06-30
;  PRIOR APPLICATION NUMBER: 60/583,621
;  PRIOR FILING DATE: 2004-06-30
;  PRIOR APPLICATION NUMBER: 60/584,829
;  PRIOR FILING DATE: 2004-06-30
;  PRIOR APPLICATION NUMBER: 60/584,800
;  PRIOR FILING DATE: 2004-06-30
;  NUMBER OF SEQ ID NOS: 2523
; SEQ ID NO 1344
;   LENGTH: 175
;   TYPE: PRT
;   ORGANISM: Triticum aestivum
;   FEATURE:
;   NAME/KEY: misc_feature
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;   LOCATION: (1)..(175)
;   OTHER INFORMATION: Public GI no. 2346976
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION:
;   OTHER INFORMATION: Utility: Useful for increasing chlorophyll and photosynthetic cap
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION:
;   OTHER INFORMATION: Utility: Useful for making ornamental plants with modified leaves
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION:
;   OTHER INFORMATION: Utility: Useful for making plants with altered leaf shape eg curl
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION:
;   OTHER INFORMATION: Utility: Useful for modifying fruit shape, composition and seed y
US-11-172-740-1344

  Query Match             47.3%;  Score 35;  DB 11;  Length 175;
  Best Local Similarity   70.0%;  Pred. No. 43;
  Matches    7;  Conservative    1;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          5 MALSKINLHN 14
              | |||:| ||
Db         25 MLLSKLNDHN 34


RESULT 13
US-11-188-298-1316
; Sequence 1316, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
;  APPLICANT: Abad, Mark S. et al.
;  TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
;  FILE REFERENCE: 38-21(53452)B
;  CURRENT APPLICATION NUMBER: US/11/188,298
;  CURRENT FILING DATE:  2005-07-22
;  PRIOR APPLICATION NUMBER: 60/592,978
;  PRIOR FILING DATE: 2004-07-31
;  NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 1316
;   LENGTH: 226
;   TYPE: PRT
;   ORGANISM: Davidia involucrata
US-11-188-298-1316

  Query Match             47.3%;  Score 35;  DB 11;  Length 226;
  Best Local Similarity   60.0%;  Pred. No. 57;
  Matches    6;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKI 10
              :|||: ||:|
Db         19 VGYGLELSRI 28


RESULT 14
US-11-188-298-5176
; Sequence 5176, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
;  APPLICANT: Abad, Mark S. et al.
;  TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
;  FILE REFERENCE: 38-21(53452)B
;  CURRENT APPLICATION NUMBER: US/11/188,298
;  CURRENT FILING DATE:  2005-07-22
;  PRIOR APPLICATION NUMBER: 60/592,978
;  PRIOR FILING DATE: 2004-07-31
;  NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 5176
;   LENGTH: 291
;   TYPE: PRT
;   ORGANISM: Glycine max
US-11-188-298-5176

  Query Match             47.3%;  Score 35;  DB 11;  Length 291;
  Best Local Similarity   60.0%;  Pred. No. 76;
  Matches    6;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKI 10
              :|||: ||:|
Db        192 IGYGLELSRI 201


RESULT 15
US-11-096-568A-20956
; Sequence 20956, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
;  APPLICANT: Alexandrov, Nickolai et al.
;  TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
;  TITLE OF INVENTION:  Therby
;  FILE REFERENCE: 2750-1592PUS2
;  CURRENT APPLICATION NUMBER: US/11/096,568A
;  CURRENT FILING DATE:  2005-04-01
;  NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 20956
;   LENGTH: 332
;   TYPE: PRT
;   ORGANISM: Zea mays subsp. mays
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)..(332)
;   OTHER INFORMATION: Ceres Seq. ID no. 12391420
US-11-096-568A-20956

  Query Match             47.3%;  Score 35;  DB 11;  Length 332;
  Best Local Similarity   30.8%;  Pred. No. 89;
  Matches    4;  Conservative    6;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              |:|:::  : :||
Db        159 GHGISIGSLGVHN 171


RESULT 16
US-11-188-298-11417
; Sequence 11417, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
;  APPLICANT: Abad, Mark S. et al.
;  TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
;  FILE REFERENCE: 38-21(53452)B
;  CURRENT APPLICATION NUMBER: US/11/188,298
;  CURRENT FILING DATE:  2005-07-22
;  PRIOR APPLICATION NUMBER: 60/592,978
;  PRIOR FILING DATE: 2004-07-31
;  NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 11417
;   LENGTH: 341
;   TYPE: PRT
;   ORGANISM: Glycine max
US-11-188-298-11417

  Query Match             47.3%;  Score 35;  DB 11;  Length 341;
  Best Local Similarity   60.0%;  Pred. No. 91;
  Matches    6;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKI 10
              :|||: ||:|
Db        192 IGYGLELSRI 201


RESULT 17
US-10-698-618-1
```

```
; Sequence 1, Application US/10698618
; Publication No. US20050271604A1
; GENERAL INFORMATION:
;  APPLICANT: GESTRELIUS, STINA
;  APPLICANT:  HAMMARSTROM, LARS
;  APPLICANT:  LYNGSTADAAS, PETTER
;  APPLICANT:  ANDERSSON, CHRISTER
;  APPLICANT:  SLABY, IVAN
;  APPLICANT:  HAMMARGREN, TOMAS
;  TITLE OF INVENTION: MATRIX PROTEIN COMPOSITIONS FOR WOUND HEALING
;  FILE REFERENCE: 47927-48292-CPA
;  CURRENT APPLICATION NUMBER: US/10/698,618
;  CURRENT FILING DATE:  2003-10-30
;  PRIOR APPLICATION NUMBER: US/10/156,300
;  PRIOR FILING DATE: 2002-05-28
;  PRIOR APPLICATION NUMBER: US/09/258,613
;  PRIOR FILING DATE: 1999-02-26
;  PRIOR APPLICATION NUMBER: DK PA 1998 01328
;  PRIOR FILING DATE: 1998-10-16
;  PRIOR APPLICATION NUMBER: 60/081,551
;  PRIOR FILING DATE: 1998-04-13
;  PRIOR APPLICATION NUMBER: DK 0270/98
;  PRIOR FILING DATE: 1998-02-27
;  NUMBER OF SEQ ID NOS: 2
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
;   LENGTH: 407
;   TYPE: PRT
;   ORGANISM: Rattus sp.
US-10-698-618-1

  Query Match             47.3%;  Score 35;  DB 9;  Length 407;
  Best Local Similarity   46.2%;  Pred. No. 1.1e+02;
  Matches    6;  Conservative    4;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLH 13
              :|:| ||: : ||
Db         70 LGFGKALNSLWLH 82


RESULT 18
US-11-096-568A-20955
; Sequence 20955, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
;  APPLICANT: Alexandrov, Nickolai et al.
;  TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
;  TITLE OF INVENTION:  Therby
;  FILE REFERENCE: 2750-1592PUS2
;  CURRENT APPLICATION NUMBER: US/11/096,568A
;  CURRENT FILING DATE:  2005-04-01
;  NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 20955
;   LENGTH: 502
;   TYPE: PRT
;   ORGANISM: Zea mays subsp. mays
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)..(502)
;   OTHER INFORMATION: Ceres Seq. ID no. 12391419
US-11-096-568A-20955

  Query Match             47.3%;  Score 35;  DB 11;  Length 502;
  Best Local Similarity   30.8%;  Pred. No. 1.4e+02;
  Matches    4;  Conservative    6;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              |:|:::  : :||
Db        329 GHGISIGSLGVHN 341


RESULT 19
US-11-096-568A-20954
; Sequence 20954, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
;  APPLICANT: Alexandrov, Nickolai et al.
;  TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
;  TITLE OF INVENTION:  Therby
;  FILE REFERENCE: 2750-1592PUS2
;  CURRENT APPLICATION NUMBER: US/11/096,568A
;  CURRENT FILING DATE:  2005-04-01
;  NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 20954
;   LENGTH: 525
;   TYPE: PRT
;   ORGANISM: Zea mays subsp. mays
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)..(525)
;   OTHER INFORMATION: Ceres Seq. ID no. 12391418
US-11-096-568A-20954

  Query Match             47.3%;  Score 35;  DB 11;  Length 525;
  Best Local Similarity   30.8%;  Pred. No. 1.5e+02;
  Matches    4;  Conservative    6;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              |:|:::  : :||
Db        352 GHGISIGSLGVHN 364


RESULT 20
US-11-188-298-7401
; Sequence 7401, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
;  APPLICANT: Abad, Mark S. et al.
;  TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
;  FILE REFERENCE: 38-21(53452)B
;  CURRENT APPLICATION NUMBER: US/11/188,298
;  CURRENT FILING DATE:  2005-07-22
;  PRIOR APPLICATION NUMBER: 60/592,978
;  PRIOR FILING DATE: 2004-07-31
;  NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 7401
;   LENGTH: 665
;   TYPE: PRT
;   ORGANISM: Glycine max
US-11-188-298-7401

  Query Match             47.3%;  Score 35;  DB 11;  Length 665;
  Best Local Similarity   60.0%;  Pred. No. 2e+02;
  Matches    6;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKI 10
              :|||: ||:|
Db        459 IGYGLELSRI 468


RESULT 21
US-11-188-298-16346
; Sequence 16346, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
;  APPLICANT: Abad, Mark S. et al.
;  TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
;  FILE REFERENCE: 38-21(53452)B
;  CURRENT APPLICATION NUMBER: US/11/188,298
;  CURRENT FILING DATE:  2005-07-22
;  PRIOR APPLICATION NUMBER: 60/592,978
;  PRIOR FILING DATE: 2004-07-31
;  NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 16346
;   LENGTH: 723
;   TYPE: PRT
```

```
;   ORGANISM: Glycine max
US-11-188-298-16346

  Query Match             47.3%;  Score 35;  DB 11;  Length 723;
  Best Local Similarity   60.0%;  Pred. No. 2.2e+02;
  Matches    6;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKI 10
              :|||: ||:|
Db        517 IGYGLELSRI 526


RESULT 22
US-11-188-298-490
; Sequence 490, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
;  APPLICANT: Abad, Mark S. et al.
;  TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
;  FILE REFERENCE: 38-21(53452)B
;  CURRENT APPLICATION NUMBER: US/11/188,298
;  CURRENT FILING DATE:  2005-07-22
;  PRIOR APPLICATION NUMBER: 60/592,978
;  PRIOR FILING DATE: 2004-07-31
;  NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 490
;   LENGTH: 724
;   TYPE: PRT
;   ORGANISM: Arabidopsis thaliana
US-11-188-298-490

  Query Match             47.3%;  Score 35;  DB 11;  Length 724;
  Best Local Similarity   60.0%;  Pred. No. 2.2e+02;
  Matches    6;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKI 10
              :|||: ||:|
Db        515 IGYGLELSRI 524


RESULT 23
US-10-995-561-526
; Sequence 526, Application US/10995561
; Publication No. US20050272054A1
; GENERAL INFORMATION:
;  APPLICANT: CARGILL, Michele et al.
;  TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
;  TITLE OF INVENTION:  CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
;  TITLE OF INVENTION:  DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001559
;  CURRENT APPLICATION NUMBER: US/10/995,561
;  CURRENT FILING DATE:  2004-11-24
;  NUMBER OF SEQ ID NOS: 85702
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 526
;   LENGTH: 1404
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-10-995-561-526

  Query Match             47.3%;  Score 35;  DB 9;  Length 1404;
  Best Local Similarity   35.7%;  Pred. No. 4.6e+02;
  Matches    5;  Conservative    6;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              :|:|   ||:::|:
Db         49 IGWGSQSSKVHIHH 62


RESULT 24
US-11-090-439-24
; Sequence 24, Application US/11090439
; Publication No. US20050266442A1
; GENERAL INFORMATION:
;  APPLICANT: Squillace, Rachel
;  APPLICANT:  Weiner, Michael P.
;  TITLE OF INVENTION: Immortalized Human Tuberous Sclerosis Null
;  TITLE OF INVENTION:  Angiomyolipoma Cell and Method of Use Thereof
;  FILE REFERENCE: 24318-502
;  CURRENT APPLICATION NUMBER: US/11/090,439
;  CURRENT FILING DATE:  2005-03-25
;  PRIOR APPLICATION NUMBER: 60/556,344
;  PRIOR FILING DATE: 2004-03-25
;  NUMBER OF SEQ ID NOS: 62
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 24
;   LENGTH: 1581
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-090-439-24

  Query Match             47.3%;  Score 35;  DB 11;  Length 1581;
  Best Local Similarity   35.7%;  Pred. No. 5.3e+02;
  Matches    5;  Conservative    6;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              :|:|   ||:::|:
Db         49 IGWGSQSSKVHIHH 62


RESULT 25
US-11-090-439-26
; Sequence 26, Application US/11090439
; Publication No. US20050266442A1
; GENERAL INFORMATION:
;  APPLICANT: Squillace, Rachel
;  APPLICANT:  Weiner, Michael P.
;  TITLE OF INVENTION: Immortalized Human Tuberous Sclerosis Null
;  TITLE OF INVENTION:  Angiomyolipoma Cell and Method of Use Thereof
;  FILE REFERENCE: 24318-502
;  CURRENT APPLICATION NUMBER: US/11/090,439
;  CURRENT FILING DATE:  2005-03-25
;  PRIOR APPLICATION NUMBER: 60/556,344
;  PRIOR FILING DATE: 2004-03-25
;  NUMBER OF SEQ ID NOS: 62
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 26
;   LENGTH: 1581
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-090-439-26

  Query Match             47.3%;  Score 35;  DB 11;  Length 1581;
  Best Local Similarity   35.7%;  Pred. No. 5.3e+02;
  Matches    5;  Conservative    6;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              :|:|   ||:::|:
Db         49 IGWGSQSSKVHIHH 62


RESULT 26
US-10-995-561-527
; Sequence 527, Application US/10995561
; Publication No. US20050272054A1
; GENERAL INFORMATION:
;  APPLICANT: CARGILL, Michele et al.
;  TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
;  TITLE OF INVENTION:  CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
;  TITLE OF INVENTION:  DETECTION AND USES THEREOF
;  FILE REFERENCE: CL001559
;  CURRENT APPLICATION NUMBER: US/10/995,561
;  CURRENT FILING DATE:  2004-11-24
;  NUMBER OF SEQ ID NOS: 85702
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 527
```

```
;   LENGTH: 1588
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-10-995-561-527

  Query Match             47.3%;  Score 35;  DB 9;  Length 1588;
  Best Local Similarity   35.7%;  Pred. No. 5.3e+02;
  Matches    5;  Conservative    6;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              :|:|   ||:::|:
Db         49 IGWGSQSSKVHIHH 62


RESULT 27
US-10-485-517-301
; Sequence 301, Application US/10485517
; Publication No. US20050256299A1
; GENERAL INFORMATION:
;  APPLICANT: University of Sheffield
;  APPLICANT:  Biosynexus Incorporated
;  APPLICANT:  Foster, Simon
;  APPLICANT:  Mond, James
;  TITLE OF INVENTION: Antigenic Polypeptides
;  FILE REFERENCE: P100629WO
;  CURRENT APPLICATION NUMBER: US/10/485,517
;  CURRENT FILING DATE:  2004-02-02
;  PRIOR APPLICATION NUMBER: GB 0118825.9
;  PRIOR FILING DATE: 2001-08-02
;  PRIOR APPLICATION NUMBER: GB 0200349.9
;  PRIOR FILING DATE: 2002-01-09
;  NUMBER OF SEQ ID NOS: 424
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 301
;   LENGTH: 174
;   TYPE: PRT
;   ORGANISM: Staphylococcus aureus
US-10-485-517-301

  Query Match             45.9%;  Score 34;  DB 9;  Length 174;
  Best Local Similarity   36.4%;  Pred. No. 65;
  Matches    4;  Conservative    6;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          3 YGMALSKINLH 13
              ||:|:: :::|
Db         81 YGLAIASLSVH 91


RESULT 28
US-11-129-143-96
; Sequence 96, Application US/11129143
; Publication No. US20050266518A1
; GENERAL INFORMATION:
;  APPLICANT: BERRY, Alan
;  APPLICANT:  BRETZEL, Werner
;  APPLICANT:  HUMBELIN, Markus
;  APPLICANT:  LOPEZ-ULIBARRI, Rual
;  APPLICANT:  MAYER, Anne F.
;  APPLICANT:  YELISEEV, Alexei A.
;  TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
;  FILE REFERENCE: C38435/121966
;  CURRENT APPLICATION NUMBER: US/11/129,143
;  CURRENT FILING DATE:  2005-05-13
;  NUMBER OF SEQ ID NOS: 197
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 96
;   LENGTH: 292
;   TYPE: PRT
;   ORGANISM: Streptococcus pyrogenes
US-11-129-143-96

  Query Match             45.9%;  Score 34;  DB 11;  Length 292;
  Best Local Similarity   66.7%;  Pred. No. 1.2e+02;
  Matches    8;  Conservative    1;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINL 12
              :||| | ||| |
Db          5 IGYGKAHSKIIL 16


RESULT 29
US-11-096-568A-29040
; Sequence 29040, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
;  APPLICANT: Alexandrov, Nickolai et al.
;  TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
;  TITLE OF INVENTION:  Therby
;  FILE REFERENCE: 2750-1592PUS2
;  CURRENT APPLICATION NUMBER: US/11/096,568A
;  CURRENT FILING DATE:  2005-04-01
;  NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 29040
;   LENGTH: 344
;   TYPE: PRT
;   ORGANISM: Arabidopsis thaliana
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)..(344)
;   OTHER INFORMATION: Ceres Seq. ID no. 3599624
US-11-096-568A-29040

  Query Match             45.9%;  Score 34;  DB 11;  Length 344;
  Best Local Similarity   38.5%;  Pred. No. 1.4e+02;
  Matches    5;  Conservative    4;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              |||:    :|:|:
Db        101 GYGIGSLTVNVHS 113


RESULT 30
US-11-096-568A-29039
; Sequence 29039, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
;  APPLICANT: Alexandrov, Nickolai et al.
;  TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
;  TITLE OF INVENTION:  Therby
;  FILE REFERENCE: 2750-1592PUS2
;  CURRENT APPLICATION NUMBER: US/11/096,568A
;  CURRENT FILING DATE:  2005-04-01
;  NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 29039
;   LENGTH: 365
;   TYPE: PRT
;   ORGANISM: Arabidopsis thaliana
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)..(365)
;   OTHER INFORMATION: Ceres Seq. ID no. 3599623
US-11-096-568A-29039

  Query Match             45.9%;  Score 34;  DB 11;  Length 365;
  Best Local Similarity   38.5%;  Pred. No. 1.5e+02;
  Matches    5;  Conservative    4;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              |||:    :|:|:
Db        122 GYGIGSLTVNVHS 134


RESULT 31
US-11-096-568A-29038
; Sequence 29038, Application US/11096568A
; Publication No. US20060048240A1
```

```
; GENERAL INFORMATION:
;  APPLICANT: Alexandrov, Nickolai et al.
;  TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
;  TITLE OF INVENTION:  Therby
;  FILE REFERENCE: 2750-1592PUS2
;  CURRENT APPLICATION NUMBER: US/11/096,568A
;  CURRENT FILING DATE:  2005-04-01
;  NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 29038
;   LENGTH: 440
;   TYPE: PRT
;   ORGANISM: Arabidopsis thaliana
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)..(440)
;   OTHER INFORMATION: Ceres Seq. ID no. 3599622
US-11-096-568A-29038

  Query Match             45.9%;  Score 34;  DB 11;  Length 440;
  Best Local Similarity   38.5%;  Pred. No. 1.9e+02;
  Matches    5;  Conservative    4;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              |||:    :|:|:
Db        197 GYGIGSLTVNVHS 209


RESULT 32
US-11-087-099-8602
; Sequence 8602, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
;  APPLICANT: Abad, Mark S. et al.
;  TITLE OF INVENTION: Genes and Uses for Plant Improvement
;  FILE REFERENCE: 38-21(53450)B EP
;  CURRENT APPLICATION NUMBER: US/11/087,099
;  CURRENT FILING DATE:  2005-03-22
;  NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 8602
;   LENGTH: 445
;   TYPE: PRT
;   ORGANISM: Butyrivibrio fibrisolvens
US-11-087-099-8602

  Query Match             45.9%;  Score 34;  DB 11;  Length 445;
  Best Local Similarity   60.0%;  Pred. No. 1.9e+02;
  Matches    6;  Conservative    2;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKIN 11
              |||:  |||:
Db        305 GYGLGYSKID 314


RESULT 33
US-11-188-298-7996
; Sequence 7996, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
;  APPLICANT: Abad, Mark S. et al.
;  TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
;  FILE REFERENCE: 38-21(53452)B
;  CURRENT APPLICATION NUMBER: US/11/188,298
;  CURRENT FILING DATE:  2005-07-22
;  PRIOR APPLICATION NUMBER: 60/592,978
;  PRIOR FILING DATE: 2004-07-31
;  NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 7996
;   LENGTH: 445
;   TYPE: PRT
;   ORGANISM: Butyrivibrio fibrisolvens
US-11-188-298-7996

  Query Match             45.9%;  Score 34;  DB 11;  Length 445;
  Best Local Similarity   60.0%;  Pred. No. 1.9e+02;
  Matches    6;  Conservative    2;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKIN 11
              |||:  |||:
Db        305 GYGLGYSKID 314


RESULT 34
US-11-079-463-7672
; Sequence 7672, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
;  APPLICANT: Gary L. Breton
;  TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES FRA
;  TITLE OF INVENTION:  FOR DIAGNOSTICS AND THERAPEUTICS
;  FILE REFERENCE: PATH00-03DIV2
;  CURRENT APPLICATION NUMBER: US/11/079,463
;  CURRENT FILING DATE:  2005-03-14
;  PRIOR APPLICATION NUMBER: US 60/128,705
;  PRIOR FILING DATE: 1999-04-09
;  PRIOR APPLICATION NUMBER: US 09/540,209
;  PRIOR FILING DATE: 2000-04-04
;  NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 7672
;   LENGTH: 467
;   TYPE: PRT
;   ORGANISM: B.fragilis
US-11-079-463-7672

  Query Match             45.9%;  Score 34;  DB 11;  Length 467;
  Best Local Similarity   66.7%;  Pred. No. 2e+02;
  Matches    6;  Conservative    2;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSK 9
              :|||: |||
Db        335 IGYGIGLSK 343


RESULT 35
US-11-079-463-8886
; Sequence 8886, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
;  APPLICANT: Gary L. Breton
;  TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES FRA
;  TITLE OF INVENTION:  FOR DIAGNOSTICS AND THERAPEUTICS
;  FILE REFERENCE: PATH00-03DIV2
;  CURRENT APPLICATION NUMBER: US/11/079,463
;  CURRENT FILING DATE:  2005-03-14
;  PRIOR APPLICATION NUMBER: US 60/128,705
;  PRIOR FILING DATE: 1999-04-09
;  PRIOR APPLICATION NUMBER: US 09/540,209
;  PRIOR FILING DATE: 2000-04-04
;  NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 8886
;   LENGTH: 521
;   TYPE: PRT
;   ORGANISM: B.fragilis
US-11-079-463-8886

  Query Match             45.9%;  Score 34;  DB 11;  Length 521;
  Best Local Similarity   70.0%;  Pred. No. 2.3e+02;
  Matches    7;  Conservative    1;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKIN 11
              || : |||||
Db        260 GYKVRLSKIN 269


RESULT 36
US-11-188-298-3229
; Sequence 3229, Application US/11188298
```

```
; Publication No. US20060075522A1
; GENERAL INFORMATION:
;  APPLICANT: Abad, Mark S. et al.
;  TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
;  FILE REFERENCE: 38-21(53452)B
;  CURRENT APPLICATION NUMBER: US/11/188,298
;  CURRENT FILING DATE:  2005-07-22
;  PRIOR APPLICATION NUMBER: 60/592,978
;  PRIOR FILING DATE: 2004-07-31
;  NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 3229
;   LENGTH: 544
;   TYPE: PRT
;   ORGANISM: Schizosaccharomyces pombe
US-11-188-298-3229

  Query Match             45.9%;  Score 34;  DB 11;  Length 544;
  Best Local Similarity   54.5%;  Pred. No. 2.4e+02;
  Matches    6;  Conservative    2;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              || |: : |||
Db        148 GMILTLVQLHN 158


RESULT 37
US-10-915-002-308
; Sequence 308, Application US/10915002
; Publication No. US20060078950A1
; GENERAL INFORMATION:
;  APPLICANT: Progulske-Fox, Ann
;  APPLICANT:  Hillman, Jeffrey D.
;  APPLICANT:  Handfield, Martin
;  TITLE OF INVENTION: IDENTIFICATION OF PORPHYROMONAS GINGIVALIS VIRULENCE POLYNUCLEOTI
;  TITLE OF INVENTION:  USE IN DIAGNOSIS ANTIGENS FOR USE IN THE DIAGNOSIS, TREATMENT, A
;  TITLE OF INVENTION:  PERIODONTAL DISEASES
;  FILE REFERENCE: 02-042
;  CURRENT APPLICATION NUMBER: US/10/915,002
;  CURRENT FILING DATE:  2004-08-10
;  NUMBER OF SEQ ID NOS: 354
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 308
;   LENGTH: 662
;   TYPE: PRT
;   ORGANISM: Porphyromonas gingivalis
US-10-915-002-308

  Query Match             45.9%;  Score 34;  DB 9;  Length 662;
  Best Local Similarity   75.0%;  Pred. No. 3e+02;
  Matches    6;  Conservative    2;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          2 GYGMALSK 9
              ||||||::
Db        327 GYGMALAQ 334


RESULT 38
US-11-072-512-2224
; Sequence 2224, Application US/11072512
; Publication No. US20060029945A1
; GENERAL INFORMATION:
;  APPLICANT: ISOGAI, TAKAO
;  APPLICANT:  SUGIYAMA, TOMOYASU
;  APPLICANT:  OTSUKI, TETSUJI
;  APPLICANT:  WAKAMATSU, AI
;  APPLICANT:  SATO, HIROYUKI
;  APPLICANT:  ISHII, SHIZUKO
;  APPLICANT:  YAMAMOTO, JUN-ICHI
;  APPLICANT:  ISONO, YUUKO
;  APPLICANT:  HIO, YURI
;  APPLICANT:  OTSUKA, KAORU
;  APPLICANT:  NAGAI, KEIICHI
;  APPLICANT:  IRIE, RYOTARO
;  APPLICANT:  TAMECHIKA, ICHIRO
;  APPLICANT:  SEKI, NAOHIKO
;  APPLICANT:  YOSHIKAWA, TSUTOMU
;  APPLICANT:  OTSUKA, MOTOYUKI
;  APPLICANT:  NAGAHARI, KENJI
;  APPLICANT:  MASUHO, YASUHIKO
;  TITLE OF INVENTION: Novel full length cDNA
;  FILE REFERENCE: 084335-0191
;  CURRENT APPLICATION NUMBER: US/11/072,512
;  CURRENT FILING DATE:  2005-03-07
;  PRIOR APPLICATION NUMBER: US 60/350,978
;  PRIOR FILING DATE: 2002-01-25
;  PRIOR APPLICATION NUMBER: JP 2001-379298
;  PRIOR FILING DATE: 2001-11-05
;  NUMBER OF SEQ ID NOS: 4096
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2224
;   LENGTH: 724
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-072-512-2224

  Query Match             45.9%;  Score 34;  DB 11;  Length 724;
  Best Local Similarity   54.5%;  Pred. No. 3.3e+02;
  Matches    6;  Conservative    3;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              | ||:::|| |
Db        493 GTALTRLNLRN 503


RESULT 39
US-11-147-047-46
; Sequence 46, Application US/11147047
; Publication No. US20050260668A1
; GENERAL INFORMATION:
;  APPLICANT: Agarwal, Pankaj
;  APPLICANT:  Murdock, Paul R.
;  APPLICANT:  Rizvi, Safia K.
;  APPLICANT:  Smith, Randall F.
;  APPLICANT:  Xiang, Zhaoying
;  TITLE OF INVENTION: NOVEL COMPOUNDS
;  FILE REFERENCE: GP50016
;  CURRENT APPLICATION NUMBER: US/11/147,047
;  CURRENT FILING DATE:  2005-06-07
;  PRIOR APPLICATION NUMBER: US/10/221,097
;  PRIOR FILING DATE: 2002-09-06
;  PRIOR APPLICATION NUMBER: PCT/US01/07143
;  PRIOR FILING DATE: 2001-03-05
;  PRIOR APPLICATION NUMBER: 60/187,107
;  PRIOR FILING DATE: 2000-03-06
;  PRIOR APPLICATION NUMBER: 60/236,874
;  PRIOR FILING DATE: 2000-10-03
;  PRIOR APPLICATION NUMBER: 60/188,916
;  PRIOR FILING DATE: 2000-03-13
;  PRIOR APPLICATION NUMBER: 60/237,846
;  PRIOR FILING DATE: 2000-10-03
;  NUMBER OF SEQ ID NOS: 52
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 46
;   LENGTH: 845
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-147-047-46

  Query Match             45.9%;  Score 34;  DB 11;  Length 845;
  Best Local Similarity   54.5%;  Pred. No. 4e+02;
  Matches    6;  Conservative    3;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              | ||:::|| |
Db        493 GTALTRLNLRN 503
```

```
RESULT 40
US-11-264-096-483
; Sequence 483, Application US/11264096
; Publication No. US20060084794A1
; GENERAL INFORMATION:
;  APPLICANT: Rosen et al.
;  TITLE OF INVENTION: Albumin Fusion Proteins
;  FILE REFERENCE: PF546D1
;  CURRENT APPLICATION NUMBER: US/11/264,096
;  CURRENT FILING DATE:  2005-11-02
;  PRIOR APPLICATION NUMBER: 09/833,245
;  PRIOR FILING DATE: 2001-04-12
;  PRIOR APPLICATION NUMBER: 60/229, 358
;  PRIOR FILING DATE: 2000-04-12
;  PRIOR APPLICATION NUMBER: 60/256, 931
;  PRIOR FILING DATE: 2000-12-21
;  PRIOR APPLICATION NUMBER: 60/199, 384
;  PRIOR FILING DATE: 2000-04-25
;  NUMBER OF SEQ ID NOS: 2267
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 483
;   LENGTH: 845
;   TYPE: PRT
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: SITE
;   LOCATION: (477)
;   OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-11-264-096-483

  Query Match             45.9%;  Score 34;  DB 11;  Length 845;
  Best Local Similarity   54.5%;  Pred. No. 4e+02;
  Matches    6;  Conservative    3;  Mismatches    2;  Indels    0;  Gaps    0;

Qy           4 GMALSKINLHN 14
               | ||:::|| |
Db         493 GTALTRLNLRN 503


Search completed: May 13, 2006, 08:29:40
Job time : 29 secs
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GenCore version 5.1.8

OM protein - protein search, using sw model

Run on: May 13, 2006, 08:10:22 ; Search time 39 Seconds
(without alignments)
34.539 Million cell updates/sec

Title: US-10-769-514-17
Perfect score: 74
Sequence: 1 MGYGMALSKINLHN 14

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 100 summaries

Database : PIR_80:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	46	62.2	188	2	AB0503	conserved hypothet
2	46	62.2	188	2	B85481	yaaH protein [simi
3	46	62.2	188	2	B90630	yaaH protein [simi
4	46	62.2	188	2	E56688	protein yaaH - Esc
5	46	62.2	203	2	AH0057	probable membrane
6	44	59.5	197	2	F82282	conserved hypothet
7	43	58.1	119	2	E72714	probable ribosomal
8	41	55.4	906	2	B96901	uncharacterized co
9	40	54.1	429	2	JC4986	site-specific DNA-
10	40	54.1	432	2	A82533	glutamyl-tRNA redu
11	40	54.1	607	2	B84153	two-component sens
12	40	54.1	1122	2	T28130	hypothetical prote
13	39.5	53.4	2338	2	I73957	kinase-related pro
14	39	52.7	89	2	A43664	usg protein - Caul
15	39	52.7	204	2	E69126	yaaH protein homol
16	39	52.7	1032	2	S74487	hypothetical prote
17	38	51.4	282	2	H97226	protein containing
18	38	51.4	316	2	S77783	hypothetical prote
19	38	51.4	444	2	T05614	hypothetical prote
20	38	51.4	583	2	T32266	hypothetical prote
21	38	51.4	627	2	T00484	hypothetical prote
22	37.5	50.7	603	2	JC7900	beta-N-acetylgluco
23	37	50.0	217	1	WJHU2C	homeotic protein H
24	37	50.0	217	1	WJMSX2	homeotic protein H
25	37	50.0	250	2	A69951	N-acetylmuramoyl-L
26	37	50.0	348	2	T40989	probable d-amino a
27	37	50.0	362	2	S27530	sporulation protei
28	37	50.0	362	2	A84963	probable proteinas
29	37	50.0	378	2	H69505	conserved hypothet
30	37	50.0	394	2	AI2206	hypothetical prote
31	37	50.0	551	2	S66740	probable transcrip
32	37	50.0	691	1	VCNVH3	capsid-associated
33	37	50.0	692	2	T41845	VP80 orf104 - Bomb
34	37	50.0	693	1	JH0265	DNA recombinase (E
35	37	50.0	693	2	G91194	DNA helicase RecG
36	37	50.0	704	2	H86041	hypothetical prote
37	37	50.0	809	2	AI2747	conserved hypothet
38	37	50.0	873	2	H97528	hypothetical prote
39	36	48.6	79	2	F72592	hypothetical prote
40	36	48.6	168	2	H83643	polypeptide deform
41	36	48.6	168	2	E87389	RNA polymerase sig
42	36	48.6	170	2	H82746	polypeptide deform
43	36	48.6	200	2	F83780	hypothetical prote
44	36	48.6	302	2	D85911	hypothetical prote
45	36	48.6	305	2	S64612	hypothetical prote
46	36	48.6	357	2	C70805	hypothetical prote
47	36	48.6	379	2	H90183	hypothetical prote
48	36	48.6	382	2	AB0945	alcohol dehydrogen
49	36	48.6	382	2	AI0040	probable methanol
50	36	48.6	409	2	S26033	NADH2 dehydrogenas
51	36	48.6	409	2	S26021	NADH2 dehydrogenas
52	36	48.6	422	2	C91067	hypothetical prote
53	36	48.6	532	2	AB1369	conserved hypothet
54	36	48.6	532	2	AC1738	conserved hypothet
55	36	48.6	573	2	B70047	two-component sens
56	36	48.6	699	2	T18984	hypothetical prote
57	36	48.6	1068	2	T48756	mitochondrial nico
58	35.5	48.0	257	1	S22363	gufA protein homol
59	35.5	48.0	257	2	AF0890	probable membrane
60	35.5	48.0	257	2	H91119	gufA protein homol
61	35.5	48.0	257	2	G85964	gufA protein homol
62	35.5	48.0	464	2	S39194	royal jelly protei
63	35.5	48.0	3036	2	T18995	hypothetical prote
64	35	47.3	132	2	T32373	hypothetical prote
65	35	47.3	207	2	S20683	phosphinomethylmal
66	35	47.3	311	2	AE3169	hypothetical prote
67	35	47.3	322	2	T27333	hypothetical prote
68	35	47.3	325	1	B40358	NADH2 dehydrogenas
69	35	47.3	325	1	DNWTU1	NADH2 dehydrogenas
70	35	47.3	325	1	S49576	NADH2 dehydrogenas
71	35	47.3	329	2	F64356	translation initia
72	35	47.3	331	1	DNOBU1	NADH2 dehydrogenas
73	35	47.3	336	2	E72359	rod shape-determin
74	35	47.3	356	2	G97072	unchracterized con
75	35	47.3	366	2	AE0105	sugar transport AT
76	35	47.3	394	2	F90598	membrane nuclease
77	35	47.3	407	2	S58062	amelin 1 - rat
78	35	47.3	413	2	AE0089	probable flagellar
79	35	47.3	498	2	T23432	hypothetical prote
80	35	47.3	499	2	E96969	histidine kinase (
81	35	47.3	514	2	S17958	cytochrome oxidase
82	35	47.3	514	2	F90770	probable third cyt
83	35	47.3	514	2	B85633	probable third cyt
84	35	47.3	567	2	C75340	probable L-lactate
85	35	47.3	569	2	H59092	hypothetical prote
86	35	47.3	580	2	C82082	penicillin-binding
87	35	47.3	598	2	T29878	hypothetical prote
88	35	47.3	631	2	G70188	transcription init
89	35	47.3	634	2	F82079	probable 2',3'-cyc
90	35	47.3	675	2	F84937	DNA ligase (NAD) (
91	35	47.3	690	1	OYRTA1	guanylate cyclase
92	35	47.3	691	1	OYBO77	guanylate cyclase
93	35	47.3	717	2	S23098	guanylate cyclase
94	35	47.3	724	2	H86427	unknown protein [i
95	35	47.3	725	2	S57127	probable membrane
96	35	47.3	732	2	S18325	guanylate cyclase,
97	35	47.3	804	2	AG1038	conserved hypothet
98	35	47.3	845	1	B71255	ribonucleoside-dip
99	35	47.3	871	2	T49216	hypothetical prote
100	35	47.3	893	2	S61000	probable membrane

ALIGNMENTS

RESULT 1
AB0503
conserved hypothetical protein STY0009 [imported] - Salmonella enterica subsp. enterica
C;Species: Salmonella enterica subsp. enterica serovar Typhi
A;Note: this species has also been called Salmonella typhi
C;Date: 09-Nov-2001 #sequence_revision 09-Nov-2001 #text_change 18-Nov-2002
C;Accession: AB0503
R;Parkhill, J.; Dougan, G.; James, K.D.; Thomson, N.R.; Pickard, D.; Wain, J.; Churcher,
th, T.; Connerton, P.; Cronin, A.; Davis, P.; Davies, R.M.; Dowd, L.; White, N.; Farrar,
, S.; Moule, S.; O'Gaora, P.
Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.;
A;Title: Complete genome sequence of a multiple drug resistant Salmonella enterica serov
A;Reference number: AB0502; MUID:21534947; PMID:11677608
A;Accession: AB0503
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-188 <PAR>
A;Cross-references: UNIPARC:UPI000005996A; GB:AL513382; PIDN:CAD01162.1; PID:g16501292;
C;Genetics:
A;Gene: STY0009
C;Superfamily: Saccharomyces cerevisiae probable membrane protein FUN34

  Query Match 62.2%; Score 46; DB 2; Length 188;
  Best Local Similarity 57.1%; Pred. No. 0.51;
  Matches 8; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
||:|| :||||
Db 15 MGFGMTTILLNLHN 28

RESULT 2
B85481
yaaH protein [similarity] - Escherichia coli (strain O157:H7, substrain EDL933)
C;Species: Escherichia coli
C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 09-Jul-2004
C;Accession: B85481
R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, D.J.; Mayhew
iller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, E.; Potamousis, K.; Apodaca,
Nature 409, 529-533, 2001
A;Title: Genome sequence of enterohemorrhagic Escherichia coli O157:H7.
A;Reference number: A85480; MUID:21074935; PMID:11206551
A;Accession: B85481
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-188 <STO>
A;Cross-references: UNIPROT:P28695; UNIPARC:UPI0000139FF9; GB:AE005174; NID:g12512685; P
A;Experimental source: strain O157:H7, substrain EDL933
C;Genetics:
A;Gene: yaaH
C;Superfamily: Saccharomyces cerevisiae probable membrane protein FUN34

  Query Match 62.2%; Score 46; DB 2; Length 188;
  Best Local Similarity 57.1%; Pred. No. 0.51;
  Matches 8; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
||:|| :||||
Db 15 MGFGMTTILLNLHN 28

RESULT 3
B90630
yaaH protein [similarity] - Escherichia coli (strain O157:H7, substrain RIMD 0509952)
C;Species: Escherichia coli
C;Date: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change 09-Jul-2004
C;Accession: B90630
R;Hayashi, T.; Makino, K.; Ohnishi, M.; Kurokawa, K.; Ishii, K.; Yokoyama, K.; Han, C.G.
gasawara, N.; Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; Shinagawa, H.
DNA Res. 8, 11-22, 2001
A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7 and geno
A;Reference number: A99629; MUID:21156231; PMID:11258796
A;Accession: B90630
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-188 <HAY>
A;Cross-references: UNIPROT:P28695; UNIPARC:UPI0000139FF9; GB:BA000007; PIDN:BAB33433.1,
A;Experimental source: strain O157:H7, substrain RIMD 0509952
C;Genetics:
A;Gene: ECs0010
C;Superfamily: Saccharomyces cerevisiae probable membrane protein FUN34

  Query Match 62.2%; Score 46; DB 2; Length 188;
  Best Local Similarity 57.1%; Pred. No. 0.51;
  Matches 8; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
||:|| :||||
Db 15 MGFGMTTILLNLHN 28

RESULT 4
E56688
protein yaaH - Escherichia coli (strain K-12)
C;Species: Escherichia coli
C;Date: 21-Jul-1995 #sequence_revision 28-Jul-1995 #text_change 09-Jul-2004
C;Accession: E56688; B64721; S28459
R;James, R.; Dean, D.O.; Debbage, J.
DNA Seq. 3, 327-332, 1993
A;Title: Five open reading frames upstream of the dnaK gene of Escherichia coli.
A;Reference number: A56688; MUID:94003405; PMID:8400364
A;Accession: E56688
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-188 <JAM>
A;Cross-references: UNIPROT:P28695; UNIPARC:UPI0000139FF9; GB:X67700; NID:g41754; PIDN:C
A;Note: authors translated the codon ATT for residue 51 as Asn
R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; Burland, V.; Riley, M.; Co
.A.; Rose, D.J.; Mau, B.; Shao, Y.
Science 277, 1453-1462, 1997
A;Title: The complete genome sequence of Escherichia coli K-12.
A;Reference number: A64720; MUID:97426617; PMID:9278503
A;Accession: B64721
A;Status: nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-188 <BLAT>
A;Cross-references: UNIPARC:UPI0000139FF9; GB:AE000111; GB:U00096; NID:g1786181; PIDN:AA
A;Experimental source: strain K-12, substrain MG1655
C;Genetics:
A;Gene: yaaH
C;Superfamily: Saccharomyces cerevisiae probable membrane protein FUN34
C;Keywords: transmembrane protein
F;37-53/Domain: transmembrane #status predicted <TM1>
F;75-91/Domain: transmembrane #status predicted <TM2>
F;98-114/Domain: transmembrane #status predicted <TM3>
F;124-140/Domain: transmembrane #status predicted <TM4>
F;149-165/Domain: transmembrane #status predicted <TM5>

  Query Match 62.2%; Score 46; DB 2; Length 188;
  Best Local Similarity 57.1%; Pred. No. 0.51;
  Matches 8; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
||:|| :||||
Db 15 MGFGMTTILLNLHN 28

RESULT 5
AH0057
probable membrane protein YPO0467 [imported] - Yersinia pestis (strain CO92)
C;Species: Yersinia pestis
C;Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 09-Jul-2004

```
C;Accession: AH0057
R;Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; Prentice, M.B.
deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.; Dougan, G.;
il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; Whitehead, S.; Barrell,
Nature 413, 523-527, 2001
A;Title: Genome sequence of Yersinia pestis, the causative agent of plague.
A;Reference number: AB0001; MUID:21470413; PMID:11586360
A;Accession: AH0057
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-203 <KUR>
A;Cross-references: UNIPROT:Q8ZIM8; UNIPARC:UPI00000DC73D; GB:AL590842; PIDN:CAC89323.1;
C;Genetics:
A;Gene: YPO0467
C;Superfamily: Saccharomyces cerevisiae probable membrane protein FUN34

  Query Match             62.2%;  Score 46;  DB 2;  Length 203;
  Best Local Similarity   57.1%;  Pred. No. 0.55;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTVLLNLHN 28


RESULT 6
F82282
conserved hypothetical protein VC0770 [imported] - Vibrio cholerae (strain N16961 serogr
C;Species: Vibrio cholerae
C;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-2004
C;Accession: F82282
R;Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, R.A.; Gwinn, M.L.; Dodson, R.J.;
chardson, D.; Ermolaeva, M.D.; Vamathevan, J.; Bass, S.; Qin, H.; Dragoi, I.; Sellers, P
l, R.R.; Mekalanos, J.J.; Venter, J.C.; Fraser, C.M.
Nature 406, 477-483, 2000
A;Title: DNA Sequence of both chromosomes of the cholera pathogen Vibrio cholerae.
A;Reference number: A82035; MUID:20406833; PMID:10952301
A;Accession: F82282
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-197 <HEI>
A;Cross-references: UNIPROT:Q9KTW0; UNIPARC:UPI00000C2DBA; GB:AE004162; GB:AE003852; NID
A;Experimental source: serogroup O1; strain N16961; biotype El Tor
C;Genetics:
A;Gene: VC0770
A;Map position: 1
C;Superfamily: Saccharomyces cerevisiae probable membrane protein FUN34

  Query Match             59.5%;  Score 44;  DB 2;  Length 197;
  Best Local Similarity   50.0%;  Pred. No. 1.3;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         14 MGFGMTTILLNIHN 27


RESULT 7
E72714
probable ribosomal protein S24 APE1132 - Aeropyrum pernix (strain K1)
C;Species: Aeropyrum pernix
C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #text_change 24-Sep-1999
C;Accession: E72714
R;Kawarabayasi, Y.; Hino, Y.; Horikawa, H.; Yamazaki, S.; Haikawa, Y.; Jin-no, K.; Takah
awa, H.; Takamiya, M.; Masuda, S.; Funahashi, T.; Tanaka, T.; Kudoh, Y.; Yamazaki, J.; K
DNA Res. 6, 83-101, 1999
A;Title: Complete genome sequence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyr
A;Reference number: A72450; MUID:99310339; PMID:10382966
A;Accession: E72714
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-119 <KAW>
A;Cross-references: UNIPARC:UPI000005DE12; DDBJ:AP000060; NID:g5104188; PIDN:BAA80117.1;
A;Experimental source: strain K1
C;Genetics:
A;Gene: APE1132

  Query Match             58.1%;  Score 43;  DB 2;  Length 119;
  Best Local Similarity   58.3%;  Pred. No. 1.1;
  Matches    7;  Conservative    2;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLH 13
              |||  |||: :|
Db         77 GYGAGLSKVRVH 88


RESULT 8
B96901
uncharacterized conserved protein, probable metal-binding CAC0010 [imported] - Clostridi
C;Species: Clostridium acetobutylicum
C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 09-Jul-2004
C;Accession: B96901
R;Nolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, K.S.; Zeng, Q.; Gibson, R.; Lee,
.; Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R.
J. Bacteriol. 183, 4823-4838, 2001
A;Title: Genome Sequence and Comparative Analysis of the Solvent-Producing Bacterium Cl
A;Reference number: A96900; MUID:21359325; PMID:21359325
A;Accession: B96901
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-906 <KUR>
A;Cross-references: UNIPROT:Q97N28; UNIPARC:UPI00001390DC; GB:AE001437; PIDN:AAK77997.1,
A;Experimental source: Clostridium acetobutylicum ATCC824
C;Genetics:
A;Gene: CAC0010

  Query Match             55.4%;  Score 41;  DB 2;  Length 906;
  Best Local Similarity   60.0%;  Pred. No. 25;
  Matches    6;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKIN 11
              |||: :||:|
Db        429 GYGLVMSKVN 438


RESULT 9
JC4986
site-specific DNA-methyltransferase (cytosine-specific) (EC 2.1.1.73) ApaLI - Acetobact
C;Species: Acetobacter pasteurianus
C;Date: 15-Dec-1996 #sequence_revision 21-Jan-1997 #text_change 31-Dec-2004
C;Accession: JC4986
R;Suzuki, T.; Sugimoto, E.; Tahara, Y.; Yamada, Y.
Biosci. Biotechnol. Biochem. 60, 1401-1405, 1996
A;Title: Cloning and nucleotide sequence of ApaLI restriction-modification system from .
A;Reference number: JC4986; MUID:97141241; PMID:8987585
A;Accession: JC4986
A;Molecule type: DNA
A;Residues: 1-429 <SUZ>
A;Cross-references: UNIPROT:P70750; UNIPARC:UPI00000B5146; DDBJ:D78276; NID:g1644233; P
A;Experimental source: strain IFO13753
C;Comment: This enzyme belongs to cytosine-5 methylase family.
C;Genetics:
A;Gene: M.ApaLI
C;Superfamily: modification methylase (cytosine-specific), M.EcoRII type
C;Keywords: methyltransferase; S-adenosylmethionine

  Query Match             54.1%;  Score 40;  DB 2;  Length 429;
  Best Local Similarity   58.3%;  Pred. No. 17;
  Matches    7;  Conservative    2;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINL 12
              ||| : | |:||
Db        142 MGYSVRLQKVNL 153


RESULT 10
```

```
A82533
glutamyl-tRNA reductase XF2648 [imported] - Xylella fastidiosa (strain 9a5c)
C;Species: Xylella fastidiosa
C;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-2004
C;Accession: A82533
R;anonymous, The Xylella fastidiosa Consortium of the Organization for Nucleotide Sequen
Nature 406, 151-157, 2000
A;Title: The genome sequence of the plant pathogen Xylella fastidiosa.
A;Reference number: A82515; MUID:20365717; PMID:10910347
A;Note: for a complete list of authors see reference number A59328 below
A;Accession: A82533
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-432 <SIM>
A;Cross-references: UNIPROT:Q9PA72; UNIPARC:UPI000012C440; GB:AE004071; GB:AE003849; NID
A;Experimental source: strain 9a5c
R;Simpson, A.J.G.; Reinach, F.C.; Arruda, P.; Abreu, F.A.; Acencio, M.; Alvarenga, R.; A
Briones, M.R.S.; Bueno, M.R.P.; Camargo, A.A.; Camargo, L.E.A.; Carraro, D.M.; Carrer, H
as-Neto, E.; Docena, C.; El-Dorry, H.; Facincani, A.P.; Ferreira, A.J.S.
submitted to GenBank, June 2000
A;Authors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.S.; Franca, S.C.; Franco, M.C.; Frohm
J.D.; Junqueira, M.L.; Kemper, E.L.; Kitajima, J.P.; Krieger, J.E.; Kuramae, E.E.; Laign
chado, M.A.; Madeira, A.M.B.N.; Madeira, H.M.F.; Marino, C.L.; Marques, M.V.; Martins, E
A;Authors: Martins, E.M.F.; Matsukuma, A.Y.; Menck, C.F.M.; Miracca, E.C.; Miyaki, C.Y.;
, F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmieri, D.A
Rodrigues, V.; Rosa, A.J. de M.; de Rosa Jr., V.E.; de Sa, R.G.; Santelli, R.V.; Sawasak
A;Authors: da Silva, A.C.R.; da Silva, F.R.; da Silva, A.M.; Silva Jr., W.A.; da Silveir
M.; Tsuhako, M.H.; Vallada, H.; Van Sluys, M.A.; Verjovski-Almeida, S.; Vettore, A.L.; Z
A;Reference number: A59328
A;Contents: annotation
C;Genetics:
A;Gene: XF2648
C;Superfamily: glutamyl-tRNA reductase

  Query Match             54.1%;  Score 40;  DB 2;  Length 432;
  Best Local Similarity   50.0%;  Pred. No. 17;
  Matches    6;  Conservative    4;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLH 13
              || : |:::|||
Db        220 GYALPLTELNLH 231


RESULT 11
B84153
two-component sensor histidine kinase BH4026 [imported] - Bacillus halodurans (strain C-
C;Species: Bacillus halodurans
C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 09-Jul-2004
C;Accession: B84153
R;Takami, H.; Nakasone, K.; Takaki, Y.; Maeno, G.; Sasaki, R.; Masui, N.; Fuji, F.; Hira
Nucleic Acids Res. 28, 4317-4331, 2000
A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans and
A;Reference number: A83650; MUID:20512582; PMID:11058132
A;Accession: B84153
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-607 <STO>
A;Cross-references: UNIPROT:Q9K5R2; UNIPARC:UPI00000C439F; GB:AP001520; GB:BA000004; NID
A;Experimental source: strain C-125
C;Genetics:
A;Gene: BH4026

  Query Match             54.1%;  Score 40;  DB 2;  Length 607;
  Best Local Similarity   50.0%;  Pred. No. 25;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||    |:|:|:
Db        221 MGYGDFSRKVNVHS 234


RESULT 12
T28130
hypothetical protein ZK970.6 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
C;Accession: T28130
R;Berks, M.
submitted to the EMBL Data Library, April 1995
A;Reference number: Z20473
A;Accession: T28130
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-1122 <WIL>
A;Cross-references: UNIPROT:Q23682; UNIPARC:UPI000007CF2C; EMBL:Z49073; PIDN:CAA88890.1,
A;Experimental source: clone ZK970
C;Genetics:
A;Gene: CESP:ZK970.6
A;Map position: 2
A;Introns: 28/3; 72/2; 153/2; 281/1; 312/3; 354/3; 401/1; 442/3; 660/3; 761/1; 819/2; 95
C;Superfamily: membrane-bound guanylate cyclase; guanylate cyclase catalytic domain homo

  Query Match             54.1%;  Score 40;  DB 2;  Length 1122;
  Best Local Similarity   77.8%;  Pred. No. 48;
  Matches    7;  Conservative    2;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          3 YGMALSKIN 11
              ||||:||:|
Db        370 YGMAVSKLN 378


RESULT 13
I73957
kinase-related protein c-ros-1 precursor - rat
N;Contains: protein-tyrosine kinase (EC 2.7.1.112) ros-1
C;Species: Rattus norvegicus (Norway rat)
C;Date: 02-Aug-1996 #sequence_revision 02-Aug-1996 #text_change 09-Jul-2004
C;Accession: I73957; I56752; I73956
R;Matsushime, H.; Shibuya, M.
J. Virol. 64, 2117-2125, 1990
A;Title: Tissue-specific expression of rat c-ros-1 gene and partial structural similarit
A;Reference number: I56752; MUID:90219211; PMID:2139140
A;Accession: I73957
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-2338 <RES>
A;Cross-references: UNIPROT:Q63132; UNIPARC:UPI00000E793D; GB:M35106; NID:g203599; PIDN
A;Accession: I56752
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-430,452-2338 <RE2>
A;Cross-references: UNIPARC:UPI00000E591B; GB:M35104; NID:g203595; PIDN:AAA40966.1; PID
A;Accession: I73956
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-430,452-1872,'AC',1875 <RE3>
A;Cross-references: UNIPARC:UPI00000E78EA; GB:M35105; NID:g203597; PIDN:AAA40967.1; PID
C;Superfamily: kinase-related protein ros; LDL receptor YWTD-containing repeat homology,
C;Keywords: alternative splicing; ATP; autophosphorylation; glycoprotein; kinase-related
ific protein kinase
F;753-793/Domain: LDL receptor YWTD-containing repeat homology <YW3>
F;1935-2214/Domain: protein kinase homology <KIN>
F;1943-1951/Region: protein kinase ATP-binding motif

  Query Match             53.4%;  Score 39.5;  DB 2;  Length 2338;
  Best Local Similarity   58.8%;  Pred. No. 1.3e+02;
  Matches   10;  Conservative    1;  Mismatches    3;  Indels    3;  Gaps    1;

Qy          1 MGY---GMALSKINLHN 14
              |||   | ||  :||||
Db       1190 MGYFAQGDALFLLNLHN 1206


RESULT 14
A43664
usg protein - Caulobacter crescentus
```

C;Species: Caulobacter crescentus
C;Date: 03-Mar-1993 #sequence_revision 03-Mar-1993 #text_change 09-Jul-2004
C;Accession: A43664; H87688
R;Ross, C.M.; Winkler, M.E.
J. Bacteriol. 170, 757-768, 1988
A;Title: Structure of the Caulobacter crescentus trpFBA operon.
A;Reference number: A43664; MUID:88115177; PMID:2828322
A;Accession: A43664
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-89 <ROS>
A;Cross-references: UNIPROT:P12288; UNIPARC:UPI0000137E4F; GB:M19129; NID:g144284; PIDN:
R;Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.E.; Eisen, J.; Heidelberg, J.
B.; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwinn, M.L.; Haft, D.H.; Kolon
n, J.; Ermolaeva, M.; White, O.; Salzberg, S.L.; Shapiro, L.; Venter, J.C.; Fraser, C.M.
Proc. Natl. Acad. Sci. U.S.A. 98, 4136-4141, 2001
A;Title: Complete Genome Sequence of Caulobacter crescentus.
A;Reference number: A87249; MUID:21173698; PMID:11259647
A;Accession: H87688
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-89 <STO>
A;Cross-references: UNIPARC:UPI0000137E4F; GB:AE005673; NID:g13425282; PIDN:AAK25508.1;
C;Genetics:
A;Gene: CC3546

  Query Match 52.7%; Score 39; DB 2; Length 89;
  Best Local Similarity 46.2%; Pred. No. 4.6;
  Matches 6; Conservative 4; Mismatches 3; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLH 13
Db 11 MGYGLTTAEIHYH 23

RESULT 15
E69126
yaaH protein homolog MTH215 - Methanobacterium thermoautotrophicum (strain Delta H)
C;Species: Methanobacterium thermoautotrophicum
C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C;Accession: E69126
R;Smith, D.R.; Doucette-Stamm, L.A.; Deloughery, C.; Lee, H.; Dubois, J.; Aldredge, T.;
; Qiu, D.; Spadafora, R.; Vicaire, R.; Wang, Y.; Wierzbowski, J.; Gibson, R.; Jiwani, N.
ki, S.; Church, G.M.; Daniels, C.J.; Mao, J.; Rice, P.; Noelling, J.; Reeve, J.N.
J. Bacteriol. 179, 7135-7155, 1997
A;Title: Complete genome sequence of Methanobacterium thermoautotrophicum Delta H: funct
A;Reference number: A69000; MUID:98037514; PMID:9371463
A;Accession: E69126
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-204 <MTH>
A;Cross-references: UNIPROT:O26317; UNIPARC:UPI000013948D; GB:AE000808; GB:AE000666; NID
A;Experimental source: strain Delta H
C;Genetics:
A;Gene: MTH215
A;Start codon: TTG
C;Superfamily: Saccharomyces cerevisiae probable membrane protein FUN34

  Query Match 52.7%; Score 39; DB 2; Length 204;
  Best Local Similarity 42.9%; Pred. No. 11;
  Matches 6; Conservative 4; Mismatches 4; Indels 0; Gaps 0;

Qy 1 MGYGMALSKINLHN 14
Db 28 LGFGITTILLNLHN 41

RESULT 16
S74487
hypothetical protein sll1060 - Synechocystis sp. (strain PCC 6803)
C;Species: Synechocystis sp.
A;Variety: PCC 6803
C;Date: 25-Apr-1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
C;Accession: S74487
R;Kaneko, T.; Sato, S.; Kotani, H.; Tanaka, A.; Asamizu, E.; Nakamura, Y.; Miyajima, N.,
o, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamada, M.; Yasuda
DNA Res. 3, 109-136, 1996
A;Title: Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis
s.
A;Reference number: S74322; MUID:97061201; PMID:8905231
A;Accession: S74487
A;Status: nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-1032 <KAN>
A;Cross-references: UNIPROT:P72637; UNIPARC:UPI0000139F8B; EMBL:D90899; GB:AB001339; NID
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1996
C;Genetics:
A;Start codon: GTG

  Query Match 52.7%; Score 39; DB 2; Length 1032;
  Best Local Similarity 63.6%; Pred. No. 68;
  Matches 7; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

Qy 2 GYGMALSKINL 12
Db 569 GYGFTLSPVNL 579

RESULT 17
H97226
protein containing uncharacterized domain from NimC family [imported] - Clostridium ace
C;Species: Clostridium acetobutylicum
C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 09-Jul-2004
C;Accession: H97226
R;Nolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, K.S.; Zeng, Q.; Gibson, R.; Lee,
.; Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R.
J. Bacteriol. 183, 4823-4838, 2001
A;Title: Genome Sequence and Comparative Analysis of the Solvent-Producing Bacterium Cl
A;Reference number: A96900; MUID:21359325; PMID:21359325
A;Accession: H97226
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-282 <KUR>
A;Cross-references: UNIPROT:Q97FS1; UNIPARC:UPI00000CA5C9; GB:AE001437; PIDN:AAK80603.1,
A;Experimental source: Clostridium acetobutylicum ATCC824
C;Genetics:
A;Gene: CAC2656

  Query Match 51.4%; Score 38; DB 2; Length 282;
  Best Local Similarity 50.0%; Pred. No. 25;
  Matches 6; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

Qy 3 YGMALSKINLHN 14
Db 231 YAIAMKKLNLNN 242

RESULT 18
S77783
hypothetical protein MC100 - Mycoplasma capricolum (fragment)
C;Species: Mycoplasma capricolum
C;Date: 09-Oct-1997 #sequence_revision 24-Oct-1997 #text_change 09-Jul-2004
C;Accession: S77783
R;Bork, P.; Ouzounis, C.; Casari, G.; Schneider, R.; Sander, C.; Dolan, M.; Gilbert, W.
Mol. Microbiol. 16, 955-967, 1995
A;Title: Exploring the Mycoplasma capricolum genome: a minimal cell reveals its physiol
A;Reference number: S77739; MUID:96059641; PMID:7476192
A;Accession: S77783
A;Status: nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-316 <BOR>
A;Cross-references: UNIPROT:Q48999; UNIPARC:UPI00000B6594; EMBL:Z33074; NID:g516148; PI
A;Experimental source: ATCC 27343
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, July 1994
C;Genetics:
A;Genetic code: SGC3

```
  Query Match             51.4%;  Score 38;  DB 2;  Length 316;
  Best Local Similarity   42.9%;  Pred. No. 29;
  Matches    6;  Conservative    5;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              :|||: |:::  ||
Db        284 IGYGVWLNRLYYHN 297


RESULT 19
T05614
hypothetical protein F9D16.290 - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 09-Jul-2004
C;Accession: T05614
R;Bevan, M.; Wedler, H.; Wedler, E.; Wambutt, R.; Hoheisel, J.; Mewes, H.W.; Mayer, K.F.
submitted to the Protein Sequence Database, February 1999
A;Reference number: Z15419
A;Accession: T05614
A;Molecule type: DNA
A;Residues: 1-444 <BEV>
A;Cross-references: UNIPROT:Q9SUP5; UNIPARC:UPI00000489D3; EMBL:AL035394
A;Experimental source: cultivar Columbia; BAC clone F9D16
C;Genetics:
A;Map position: 4
A;Introns: 110/3; 151/1; 209/2; 233/1
A;Note: F9D16.290
C;Superfamily: polygalacturonase

  Query Match             51.4%;  Score 38;  DB 2;  Length 444;
  Best Local Similarity   70.0%;  Pred. No. 41;
  Matches    7;  Conservative    2;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLH 13
              |: ||:|||| 
Db        393 GICLSEINLH 402


RESULT 20
T32266
hypothetical protein F23F1.6 - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change 09-Jul-2004
C;Accession: T32266
R;Wu, X.
submitted to the EMBL Data Library, September 1997
A;Description: The sequence of C. elegans cosmid F23F1.
A;Reference number: Z21142
A;Accession: T32266
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-583 <WUX>
A;Cross-references: UNIPROT:O17069; UNIPARC:UPI000007E396; EMBL:AF024493; PIDN:AAB70324.
A;Experimental source: strain Bristol N2; clone F23F1
C;Genetics:
A;Gene: CESP:F23F1.6
A;Map position: 2
A;Introns: 88/2; 155/3; 199/3; 428/1; 509/3
C;Superfamily: ecotropic retrovirus receptor protein

  Query Match             51.4%;  Score 38;  DB 2;  Length 583;
  Best Local Similarity   63.6%;  Pred. No. 56;
  Matches    7;  Conservative    2;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |  ||||::||
Db        561 GQKLSKIDVHN 571


RESULT 21
T00484
hypothetical protein At2g35030 [imported] - Arabidopsis thaliana
N;Alternate names: hypothetical protein F19I3.26
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 12-Feb-1999 #sequence_revision 12-Feb-1999 #text_change 09-Jul-2004
C;Accession: T00484; G84763
R;Rounsley, S.D.; Lin, X.; Ketchum, K.A.; Crosby, M.L.; Brandon, R.C.; Sykes, S.M.; Kaul
submitted to the EMBL Data Library, April 1998
A;Description: Arabidopsis thaliana chromosome II BAC F19I3 genomic sequence.
A;Reference number: Z14160
A;Accession: T00484
A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-627 <ROU>
A;Cross-references: UNIPROT:O64766; UNIPARC:UPI00000A16AD; EMBL:AC004238; NID:g3033373;
A;Experimental source: cultivar Columbia
R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y.;
M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon, L
euss, D.; Nierman, W.C.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter, J
Nature 402, 761-768, 1999
A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana.
A;Reference number: A84420; MUID:20083487; PMID:10617197
A;Accession: G84763
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-627 <STO>
A;Cross-references: UNIPARC:UPI00000A16AD; GB:AE002093; NID:g3033399; PIDN:AAC12843.1; G
C;Genetics:
A;Gene: F19I3.26; At2g35030
A;Map position: 2

  Query Match             51.4%;  Score 38;  DB 2;  Length 627;
  Best Local Similarity   58.3%;  Pred. No. 61;
  Matches    7;  Conservative    1;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          3 YGMALSKINLHN 14
              ||  ||  |:||
Db        505 YGAILSACNVHN 516


RESULT 22
JC7900
beta-N-acetylglucosaminidase (EC 3.2.1.30) NagA - Emericella nidulans, Aspergillus nidul
C;Species: Emericella nidulans, Aspergillus nidulans
C;Date: 03-Feb-2003 #sequence_revision 03-Feb-2003 #text_change 09-Jul-2004
C;Accession: JC7900
R;Kim, S.; Matsuo, I.; Ajisaka, K.; Nakajima, H.; Kitamoto, K.
Biosci. Biotechnol. Biochem. 66, 2168-2175, 2002
A;Title: Cloning and characterization of the nagA gene that encodes beta-N-acetylglucosa
A;Reference number: JC7900; MUID:22333931; PMID:12450128
A;Accession: JC7900
A;Molecule type: mRNA
A;Residues: 1-603 <KIM>
A;Cross-references: UNIPROT:Q9HGI3; UNIPARC:UPI0000069AFD; DDBJ:AB039846
C;Comment: This enzyme is generally dimeric and has broad substrate specificity. It has
gradation of chitin cell wall by endochitinases.
C;Genetics:
A;Gene: nagA
C;Keywords: glycosidase; hydrolase

  Query Match             50.7%;  Score 37.5;  DB 2;  Length 603;
  Best Local Similarity   81.8%;  Pred. No. 72;
  Matches    9;  Conservative    1;  Mismatches    0;  Indels    1;  Gaps    1;

Qy          4 GMALSKIN-LH 13
              ||||||:| ||
Db        210 GMALSKLNVLH 220


RESULT 23
WJHU2C
homeotic protein Hox B7 - human
N;Alternate names: homeotic protein c1; homeotic protein Hox 2C; TATAA binding protein
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 09-Jul-2004
```

C;Accession: A28030; S15535; A44934
R;Simeone, A.; Mavilio, F.; Acampora, D.; Giampaolo, A.; Faiella, A.; Zappavigna, V.; D'
Proc. Natl. Acad. Sci. U.S.A. 84, 4914-4918, 1987
A;Title: Two human homeobox genes, c1 and c8: structure analysis and expression in embry
A;Reference number: A28030; MUID:87260899; PMID:2885844
A;Accession: A28030
A;Molecule type: mRNA
A;Residues: 1-217 <SIM>
A;Cross-references: UNIPROT:P09629; UNIPARC:UPI00001745A9; GB:M16937
A;Note: the authors translated the codon GGC for residue 53 as Ala
R;Boncinelli, E.; Acampora, D.; Pannese, M.; d'Esposito, M.; Somma, R.; Gaudino, G.; Sto
Genome 31, 745-756, 1989
A;Title: Organization of human class I homeobox genes.
A;Reference number: S15036; MUID:90215256; PMID:2576652
A;Accession: S15535
A;Molecule type: DNA
A;Residues: 137-202 <BON>
A;Cross-references: UNIPARC:UPI00001745AA
R;Baier, L.J.; Hannibal, M.C.; Hanley, E.W.; Nabel, G.J.
Blood 78, 1047-1055, 1991
A;Title: Lymphoid expression and TATAA binding of a human protein containing an Antennap
A;Reference number: A44934; MUID:91329816; PMID:1678287
A;Accession: A44934
A;Molecule type: mRNA
A;Residues: 98-217 <BAI>
A;Cross-references: UNIPARC:UPI00001745AB; GB:S49765; NID:g233572; PIDN:AAB19469.1; PID:
A;Note: this sequence is inconsistent with the nucleotide translation
A;Note: sequence extracted from NCBI backbone (NCBIN:49765, NCBIP:49769)
C;Genetics:
A;Gene: GDB:HOXB7
A;Cross-references: GDB:120660; OMIM:142962
A;Map position: 17q21.3-17q21.3
A;Introns: 134/1
C;Superfamily: homeotic protein Hox A7; homeobox homology
C;Keywords: DNA binding; homeobox; nucleus; transcription regulation
F;138-194/Domain: homeobox homology <HOX>

  Query Match 50.0%; Score 37; DB 1; Length 217;
  Best Local Similarity 50.0%; Pred. No. 29;
  Matches 6; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

Qy 2 GYGMALSKINLH 13
|||: | |:|
Db 77 GYGLEPSSFNMH 88

RESULT 24
WJMSX2
homeotic protein Hox B7 - mouse
N;Alternate names: homeotic protein Hox 2.3
C;Species: Mus musculus (house mouse)
C;Date: 30-Jun-1991 #sequence_revision 30-Jun-1991 #text_change 09-Jul-2004
C;Accession: A26846; B26846; B27176; A29585; S00988; I48411; S01887
R;Meijlink, F.; de Laaf, R.; Verrijzer, P.; Destree, O.; Kroezen, V.; Hilkens, J.; Desch
Nucleic Acids Res. 15, 6773-6786, 1987
A;Title: A mouse homeobox containing gene on chromosome 11: sequence and tissue-specific
A;Reference number: A26846; MUID:88015526; PMID:2889183
A;Accession: A26846
A;Molecule type: DNA
A;Residues: 1-217 <MEI>
A;Cross-references: UNIPROT:P09024; UNIPARC:UPI0000029981; GB:Y00436; NID:g51387; PIDN:C
A;Accession: B26846
A;Molecule type: mRNA
A;Residues: 1-217 <ME2>
A;Cross-references: UNIPARC:UPI0000029981; EMBL:Y00436; NID:g51387; PIDN:CAA68494.1; PID
R;Hart, C.P.; Fainsod, A.; Ruddle, F.H.
Genomics 1, 182-195, 1987
A;Title: Sequence analysis of the murine Hox-2.2, -2.3, and -2.4 homeo boxes: evolutiona
A;Reference number: A27176; MUID:88085193; PMID:2891608
A;Accession: B27176
A;Molecule type: DNA
A;Residues: 134-210,'R',212-217 <HAR>
A;Cross-references: UNIPARC:UPI00001745AC; EMBL:M18400
R;Lonai, P.; Arman, E.; Czosnek, H.; Ruddle, F.H.; Blatt, C.
DNA 6, 409-418, 1987
A;Title: New murine homeoboxes: structure, chromosomal assignment, and differential expr
A;Reference number: A29585; MUID:88054465; PMID:2890503
A;Accession: A29585
A;Molecule type: DNA
A;Residues: 'LCV',134-185,'G',187-205,'H',207-210,'A',212-217 <LON>
A;Cross-references: UNIPARC:UPI00001745AD; EMBL:M18167
A;Note: the authors translated the codon CAG for residue 186 as Gly
R;Kongsuwan, K.; Webb, E.; Housiaux, P.; Adams, J.M.
EMBO J. 7, 2131-2138, 1988
A;Title: Expression of multiple homeobox genes within diverse mammalian haemopoietic lir
A;Reference number: S00987; MUID:88329001; PMID:2901346
A;Accession: S00988
A;Molecule type: mRNA
A;Residues: 137-196 <KON>
A;Cross-references: UNIPARC:UPI000016CDEF; EMBL:X14570; NID:g51388; PIDN:CAA32708.1; PI
R;Verrijzer, P.; de Graaff, W.; Deschamps, J.; Meijlink, F.
Nucleic Acids Res. 16, 2729, 1988
A;Title: Nucleotide sequence of the Hox2.3 gene region.
A;Reference number: I48411; MUID:88203221; PMID:2896332
A;Accession: I48411
A;Status: translation not shown; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-217 <RES>
A;Cross-references: UNIPARC:UPI0000029981; EMBL:X06762; NID:g51389; PIDN:CAA29934.1; PI
C;Genetics:
A;Gene: Hox-2.3
A;Map position: 11
C;Superfamily: homeotic protein Hox A7; homeobox homology
C;Keywords: DNA binding; homeobox; nucleus; transcription regulation
F;138-194/Domain: homeobox homology <HOX>

  Query Match 50.0%; Score 37; DB 1; Length 217;
  Best Local Similarity 50.0%; Pred. No. 29;
  Matches 6; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

Qy 2 GYGMALSKINLH 13
|||: | |:|
Db 77 GYGLEPSSFNMH 88

RESULT 25
A69951
N-acetylmuramoyl-L-alanine amidase homolog yqeE - Bacillus subtilis
C;Species: Bacillus subtilis
C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C;Accession: A69951
R;Kunst, F.; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Berte
C.; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Ch
A.; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
Nature 390, 249-256, 1997
A;Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Galle
iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F
Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois
A;Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Mauee
y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetell
Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon
A;Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Sero
akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchiyama
T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida,
A;Authors: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A.
A;Title: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis.
A;Reference number: A69580; MUID:98044033; PMID:9384377
A;Accession: A69951
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-250 <KUN>
A;Cross-references: UNIPROT:P54450; UNIPARC:UPI000006076F; GB:Z99117; GB:AL009126; NID:
A;Experimental source: strain 168
C;Genetics:
A;Gene: yqeE
C;Superfamily: Bacillus N-acetylmuramoyl-L-alanine amidase

```
  Query Match             50.0%;  Score 37;  DB 2;  Length 250;
  Best Local Similarity   53.8%;  Pred. No. 34;
  Matches    7;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              || || : | :||
Db         18 GYAMAPAYITVHN 30


RESULT 26
T40989
probable d-amino acid oxidase - fission yeast (Schizosaccharomyces pombe)
C;Species: Schizosaccharomyces pombe
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
C;Accession: T40989
R;Lyne, M.; Rajandream, M.A.; Barrell, B.G.; Volckaert, G.
submitted to the EMBL Data Library, March 1999
A;Reference number: Z21962
A;Accession: T40989
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-348 <LYN>
A;Cross-references: UNIPROT:Q9Y7N4; UNIPARC:UPI000006B2DF; EMBL:AL049559; PIDN:CAB40174.
A;Experimental source: strain 972h-; cosmid c1450
C;Genetics:
A;Gene: SPDB:SPCC1450.07c
A;Map position: 3
C;Superfamily: D-amino-acid oxidase

  Query Match             50.0%;  Score 37;  DB 2;  Length 348;
  Best Local Similarity   63.6%;  Pred. No. 49;
  Matches    7;  Conservative    1;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINL 12
              ||||||  : |
Db        330 GYGMALDSVML 340


RESULT 27
S27530
sporulation protein - Clostridium acetobutylicum
C;Species: Clostridium acetobutylicum
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004
C;Accession: S27530
R;Reid, S.J.; Hancock, K.; Santangelo, J.D.; Woods, D.R.
submitted to the EMBL Data Library, March 1992
A;Description: Cloning and sequencing of a spoIID gene from Clostridium acetobutylicum.
A;Reference number: S27530
A;Accession: S27530
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-362 <REI>
A;Cross-references: UNIPROT:Q45833; UNIPARC:UPI00000B7749; EMBL:M87835; NID:g144914; PID
C;Superfamily: stage II sporulation protein D

  Query Match             50.0%;  Score 37;  DB 2;  Length 362;
  Best Local Similarity   53.8%;  Pred. No. 51;
  Matches    7;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLHN 14
              | | ::|| || |
Db         35 GSGFSISKFNLDN 47


RESULT 28
A84963
probable proteinase sohB [imported] - Buchnera sp. (strain APS)
C;Species: Buchnera sp.
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 31-Dec-2004
C;Accession: A84963
R;Shigenobu, S.; Watanabe, H.; Hattori, M.; Sakaki, Y.; Ishikawa, H.
Nature 407, 81-86, 2000
A;Title: Genome sequence of the endocellular bacterial symbiont of aphids Buchnera sp. A
A;Reference number: A84930; MUID:20445173; PMID:10993077
A;Accession: A84963
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-362 <STO>
A;Cross-references: UNIPARC:UPI000005E514; GB:AP000398; GSPDB:GN00144
A;Experimental source: strain APS
C;Genetics:
A;Gene: sohB; BU283
C;Superfamily: short protease IV-related protein

  Query Match             50.0%;  Score 37;  DB 2;  Length 362;
  Best Local Similarity   60.0%;  Pred. No. 51;
  Matches    6;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKIN 11
              |||:| |::|
Db        165 GYGLAASQLN 174


RESULT 29
H69505
conserved hypothetical protein AF2049 - Archaeoglobus fulgidus
C;Species: Archaeoglobus fulgidus
C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C;Accession: H69505
R;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Dodson
.; Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, E.F
 Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; McDonald, L.
Nature 390, 364-370, 1997
A;Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; Sykes, S
Smith, H.O.; Woese, C.R.; Venter, J.C.
A;Title: The complete genome sequence of the hyperthermophilic, sulfate-reducing archaeo
A;Reference number: A69250; MUID:98049343; PMID:9389475
A;Accession: H69505
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-378 <KLE>
A;Cross-references: UNIPROT:O28230; UNIPARC:UPI0000056AAD; GB:AE000961; GB:AE000782; NID

  Query Match             50.0%;  Score 37;  DB 2;  Length 378;
  Best Local Similarity   58.3%;  Pred. No. 54;
  Matches    7;  Conservative    2;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINL 12
              |||| | : :||
Db        118 MGYGFARNCVNL 129


RESULT 30
AI2206
hypothetical protein all3208 [imported] - Nostoc sp. (strain PCC 7120)
C;Species: Nostoc sp. PCC 7120
A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. strain PCC 7120
C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change 09-Jul-2004
C;Accession: AI2206
R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, A.; Iriguchi
 Nakazaki, N.; Shimpo, S.; Sugimoto, M.; Takazawa, M.; Yamada, M.; Yasuda, M.; Tabata, S
DNA Res. 8, 205-213, 2001
A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium Ana
A;Reference number: AB1807; MUID:21595285; PMID:11759840
A;Accession: AI2206
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-394 <KUR>
A;Cross-references: UNIPROT:Q8YS83; UNIPARC:UPI00000CE6E4; GB:BA000019; PIDN:BAB74907.1,
A;Experimental source: strain PCC 7120
C;Genetics:
A;Gene: all3208
C;Superfamily: Synechocystis hypothetical protein slr1087

  Query Match             50.0%;  Score 37;  DB 2;  Length 394;
```

```
  Best Local Similarity   50.0%;  Pred. No. 56;
  Matches    6;  Conservative    5;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINL 12
              :|:|:|| |::|
Db        286 LGFGVALFKVSL 297


RESULT 31
S66740
probable transcription factor YOL055c - yeast (Saccharomyces cerevisiae)
N;Alternate names: protein O1239
C;Species: Saccharomyces cerevisiae
C;Date: 12-Jul-1996 #sequence_revision 12-Jul-1996 #text_change 09-Jul-2004
C;Accession: S66740; S66747; S59294; S61724
R;Ansorge, W.; Benes, V.; Rechmann, S.; Schwager, C.; Teodoru, C.; Voss, H.; Wiemann, S.
submitted to the Protein Sequence Database, July 1996
A;Reference number: S66723
A;Accession: S66740
A;Molecule type: DNA
A;Residues: 1-551 <ANS>
A;Cross-references: UNIPROT:Q08224; UNIPARC:UPI000006B390; EMBL:Z74797; NID:g1419864; PI
A;Experimental source: strain S288C
R;Feldmann, H.; Mannhaupt, G.; Vetter, I.
submitted to the Protein Sequence Database, July 1996
A;Reference number: S66743
A;Accession: S66747
A;Molecule type: DNA
A;Residues: 1-551 <FEL>
A;Cross-references: UNIPARC:UPI000006B390; EMBL:Z74797; NID:g1419864; PID:e251864; PID:g
A;Experimental source: strain S288C
R;Mannhaupt, G.; Vetter, I.; Schwarzlose, C.; Mitzel, S.; Feldmann, H.
submitted to the EMBL Data Library, August 1995
A;Description: Analysis of a 26kb region on the left arm of yeast chromosome XV.
A;Reference number: S59285
A;Accession: S59294
A;Molecule type: DNA
A;Residues: 1-543 <FEW>
A;Cross-references: UNIPARC:UPI000006ACE0; EMBL:X91067; NID:g984177; PID:g984187
R;Mannhaupt, G.; Vetter, I.; Schwarzlose, C.; Mitzel, S.; Feldmann, H.
Yeast 12, 67-76, 1996
A;Title: Analysis of a 26 kb region on the left arm of yeast chromosome XV.
A;Reference number: S61715; MUID:96381248; PMID:8789261
A;Accession: S61724
A;Status: nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-543 <MAN>
A;Cross-references: UNIPARC:UPI000006ACE0; EMBL:X91067; NID:g984177; PIDN:CAA62531.1; PI
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, August 1995
C;Genetics:
A;Gene: SGD:THI20
A;Cross-references: SGD:S0005416
A;Map position: 15L

  Query Match             50.0%;  Score 37;  DB 2;  Length 551;
  Best Local Similarity   70.0%;  Pred. No. 81;
  Matches    7;  Conservative    2;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKI 10
              |||| ||:|:
Db        470 MGYGEALTKM 479


RESULT 32
VCNVH3
capsid-associated protein - Autographa californica nuclear polyhedrosis virus
C;Species: Autographa californica nuclear polyhedrosis virus, AcMNPV
C;Date: 30-Jun-1993 #sequence_revision 30-Jun-1993 #text_change 09-Jul-2004
C;Accession: A43376; A72863; S27897
R;Lu, A.; Carstens, E.B.
Virology 190, 201-209, 1992
A;Title: Nucleotide sequence and transcriptional analysis of the p80 gene of Autographa
id-associated gene.
A;Reference number: A43376; MUID:92410596; PMID:1529529
A;Accession: A43376
A;Molecule type: DNA
A;Residues: 1-691 <LUA>
A;Cross-references: UNIPROT:Q00733; UNIPARC:UPI0000138C95; GB:M94914; NID:g332467; PIDN
A;Experimental source: strain HR3
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1992
R;Ayres, M.D.; Howard, S.C.; Kuzio, J.; Lopez-Ferber, M.; Possee, R.D.
Virology 202, 586-605, 1994
A;Title: The complete DNA sequence of Autographa californica nuclear polyhedrosis virus
A;Reference number: A72850; MUID:94303173; PMID:8030224
A;Accession: A72863
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-691 <AYR>
A;Cross-references: UNIPARC:UPI0000138C95; GB:L22858; NID:g510708; PIDN:AAA66734.1; PID
C;Genetics:
A;Gene: p80; Ac-vp80
A;Map position: 67.2-68.5
C;Superfamily: baculovirus p87 capsid protein
C;Keywords: capsid protein; glycoprotein
F;2,71,102,319/Binding site: carbohydrate (Asn) (covalent) #status predicted

  Query Match             50.0%;  Score 37;  DB 1;  Length 691;
  Best Local Similarity   50.0%;  Pred. No. 1e+02;
  Matches    6;  Conservative    3;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          3 YGMALSKINLHN 14
              ||  | ::||:|
Db        527 YGSLLKRLNLYN 538


RESULT 33
T41845
VP80 orf104 - Bombyx mori nuclear polyhedrosis virus (isolate T3)
C;Species: Bombyx mori nuclear polyhedrosis virus, BmSNPV
A;Variety: isolate T3
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
C;Accession: T41845
R;Gomi, S.; Majima, K.; Maeda, S.
J. Gen. Virol. 80, 1323-1337, 1999
A;Title: Sequence analysis of the genome of Bombyx mori nucleopolyhedrovirus.
A;Reference number: Z22020; MUID:99281911; PMID:10355780
A;Accession: T41845
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-692 <KAM>
A;Cross-references: UNIPROT:O92464; UNIPARC:UPI00000F9A16; EMBL:L33180; NID:g3745835; P
A;Experimental source: isolate T3
C;Genetics:
A;Note: vp80
C;Superfamily: baculovirus p87 capsid protein

  Query Match             50.0%;  Score 37;  DB 2;  Length 692;
  Best Local Similarity   50.0%;  Pred. No. 1e+02;
  Matches    6;  Conservative    3;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          3 YGMALSKINLHN 14
              ||  | ::||:|
Db        528 YGSLLKRLNLYN 539


RESULT 34
JH0265
DNA recombinase (EC 3.6.1.-) - Escherichia coli (strain K-12)
N;Alternate names: RecG protein
C;Species: Escherichia coli
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 09-Jul-2004
C;Accession: JH0265; S18195; F65166
R;Kalman, M.; Murphy, H.; Cashel, M.
Gene 110, 95-99, 1992
A;Title: The nucleotide sequence of recG, the distal spo operon gene in Escherichia col
A;Reference number: JH0265; MUID:92184121; PMID:1544582
```

A;Accession: JH0265
A;Molecule type: DNA
A;Residues: 1-693 <KAL>
A;Cross-references: UNIPROT:P24230; UNIPARC:UPI000003EB2A; GB:M64367; NID:g147543; PIDN:
A;Experimental source: strain K12
R;Lloyd, R.G.; Sharples, G.J.
J. Bacteriol. 173, 6837-6843, 1991
A;Title: Molecular organization and nucleotide sequence of the recG locus of Escherichia
A;Reference number: S18195; MUID:92041567; PMID:1938888
A;Accession: S18195
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-693 <LLO>
A;Cross-references: UNIPARC:UPI000003EB2A; EMBL:X59550; NID:g42668; PIDN:CAA42123.1; PID
R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; Burland, V.; Riley, M.; Co
.A.; Rose, D.J.; Mau, B.; Shao, Y.
Science 277, 1453-1462, 1997
A;Title: The complete genome sequence of Escherichia coli K-12.
A;Reference number: A64720; MUID:97426617; PMID:9278503
A;Accession: F65166
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-693 <BLAT>
A;Cross-references: UNIPARC:UPI000003EB2A; GB:AE000442; GB:U00096; NID:g2367253; PIDN:AA
A;Experimental source: strain K-12, substrain MG1655
C;Genetics:
A;Gene: recG
A;Map position: 82 min
C;Superfamily: DNA helicase recG
C;Keywords: ATP; DNA binding; hydrolase; nucleotide binding; P-loop
F;296-303/Region: nucleotide-binding motif A (P-loop)
F;393-398/Region: nucleotide-binding motif B
F;397-400/Region: DEXH motif

  Query Match 50.0%; Score 37; DB 1; Length 693;
  Best Local Similarity 62.5%; Pred. No. 1e+02;
  Matches 10; Conservative 0; Mismatches 2; Indels 4; Gaps 1;

Qy 2 GYGMALS----KINLH 13
 | | ||| |||||
Db 16 GVGAALSNKLAKINLH 31

RESULT 35
G91194
DNA helicase RecG [imported] - Escherichia coli (strain O157:H7, substrain RIMD 0509952)
C;Species: Escherichia coli
C;Date: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change 09-Jul-2004
C;Accession: G91194
R;Hayashi, T.; Makino, K.; Ohnishi, M.; Kurokawa, K.; Ishii, K.; Yokoyama, K.; Han, C.G.
gasawara, N.; Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; Shinagawa, H.
DNA Res. 8, 11-22, 2001
A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7 and genc
A;Reference number: A99629; MUID:21156231; PMID:11258796
A;Accession: G91194
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-693 <HAY>
A;Cross-references: UNIPROT:Q8XD86; UNIPARC:UPI0000133577; GB:BA000007; PIDN:BAB37950.1;
A;Experimental source: strain O157:H7, substrain RIMD 0509952
C;Genetics:
A;Gene: ECs4527
C;Superfamily: DNA helicase recG

  Query Match 50.0%; Score 37; DB 2; Length 693;
  Best Local Similarity 62.5%; Pred. No. 1e+02;
  Matches 10; Conservative 0; Mismatches 2; Indels 4; Gaps 1;

Qy 2 GYGMALS----KINLH 13
 | | ||| |||||
Db 16 GVGAALSNKLAKINLH 31

RESULT 36
H86041
hypothetical protein recG [imported] - Escherichia coli (strain O157:H7, substrain EDL93
C;Species: Escherichia coli
C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 14-Sep-2001
C;Accession: H86041
R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, D.J.; Mayhew
iller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, E.; Potamousis, K.; Apodaca,
Nature 409, 529-533, 2001
A;Title: Genome sequence of enterohemorrhagic Escherichia coli O157:H7.
A;Reference number: A85480; MUID:21074935; PMID:11206551
A;Accession: H86041
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-704 <STO>
A;Cross-references: UNIPARC:UPI000016597F; GB:AE005174; NID:g12518411; PIDN:AAG58796.1,
A;Experimental source: strain O157:H7, substrain EDL933
C;Genetics:
A;Gene: recG
C;Superfamily: DNA helicase recG

  Query Match 50.0%; Score 37; DB 2; Length 704;
  Best Local Similarity 62.5%; Pred. No. 1.1e+02;
  Matches 10; Conservative 0; Mismatches 2; Indels 4; Gaps 1;

Qy 2 GYGMALS----KINLH 13
 | | ||| |||||
Db 27 GVGAALSNKLAKINLH 42

RESULT 37
AI2747
conserved hypothetical protein Atu1393 [imported] - Agrobacterium tumefaciens (strain C5
C;Species: Agrobacterium tumefaciens
C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 09-Jul-2004
C;Accession: AI2747
R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, Y.; Woo, L
erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, M.; McClell
; Karp, P.; Romero, P.; Zhang, S.
Science 294, 2317-2323, 2001
A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm,
ster, E.W.
A;Title: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens C58.
A;Reference number: AB2577; MUID:21608550; PMID:11743193
A;Accession: AI2747
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-809 <KUR>
A;Cross-references: UNIPROT:Q8UFK4; UNIPARC:UPI00001645E0; GB:AE008688; PIDN:AAL42399.1,
A;Experimental source: strain C58 (Dupont)
C;Genetics:
A;Gene: Atu1393
A;Map position: circular chromosome

  Query Match 50.0%; Score 37; DB 2; Length 809;
  Best Local Similarity 100.0%; Pred. No. 1.2e+02;
  Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MGYGMAL 7
 |||||||
Db 524 MGYGMAL 530

RESULT 38
H97528
hypothetical protein AGR_C_2573 [imported] - Agrobacterium tumefaciens (strain C58, Cere
C;Species: Agrobacterium tumefaciens
C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 09-Jul-2004
C;Accession: H97528
R;Goodner, B.; Hinkle, G.; Gattung, S.; Miller, N.; Blanchard, M.; Qurollo, B.; Goldman,
A.; Liu, F.; Wollam, C.; Allinger, M.; Doughty, D.; Scott, C.; Lappas, C.; Markelz, B ,
Science 294, 2323-2328, 2001
A;Title: Genome Sequence of the Plant Pathogen and Biotechnology Agent Agrobacterium tum

```
A;Reference number: A97359; MUID:21608551; PMID:11743194
A;Accession: H97528
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-873 <KUR>
A;Cross-references: UNIPROT:Q8UFK4; UNIPARC:UPI00000D1B38; GB:AE007869; PIDN:AAK87185.1;
C;Genetics:
A;Gene: AGR_C_2573
A;Map position: circular chromosome

  Query Match             50.0%;  Score 37;  DB 2;  Length 873;
  Best Local Similarity   100.0%;  Pred. No. 1.3e+02;
  Matches    7;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMAL 7
              |||||||
Db        588 MGYGMAL 594


RESULT 39
F72592
hypothetical protein APES043 - Aeropyrum pernix (strain K1)
C;Species: Aeropyrum pernix
C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #text_change 09-Jul-2004
C;Accession: F72592
R;Kawarabayasi, Y.; Hino, Y.; Horikawa, H.; Yamazaki, S.; Haikawa, Y.; Jin-no, K.; Takah
awa, H.; Takamiya, M.; Masuda, S.; Funahashi, T.; Tanaka, T.; Kudoh, Y.; Yamazaki, J.; K
DNA Res. 6, 83-101, 1999
A;Title: Complete genome sequence of an aerobic hyper-thermophilic Crenarchaeon, Aeropyr
A;Reference number: A72450; MUID:99310339; PMID:10382966
A;Accession: F72592
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-79 <KAW>
A;Cross-references: UNIPROT:Q9YCQ1; UNIPARC:UPI000005DE61; DDBJ:AP000061; NID:g5104821;
A;Experimental source: strain K1
C;Genetics:
A;Gene: APES043

  Query Match             48.6%;  Score 36;  DB 2;  Length 79;
  Best Local Similarity   60.0%;  Pred. No. 15;
  Matches    6;  Conservative    3;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKI 10
              ||:| |:||:
Db          1 MGWGCAMSKV 10


RESULT 40
H83643
polypeptide deformylase PA0019 [imported] - Pseudomonas aeruginosa (strain PAO1)
C;Species: Pseudomonas aeruginosa
C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004
C;Accession: H83643
R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, M.J.; Br
adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, K.; Lim,
.; Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic patho
A;Reference number: A82950; MUID:20437337; PMID:10984043
A;Accession: H83643
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-168 <STO>
A;Cross-references: UNIPROT:Q9I7A8; UNIPARC:UPI000012915C; GB:AE004441; GB:AE004091; NID
A;Experimental source: strain PAO1
C;Genetics:
A;Gene: def; PA0019
C;Superfamily: peptide deformylase

  Query Match             48.6%;  Score 36;  DB 2;  Length 168;
  Best Local Similarity   41.7%;  Pred. No. 34;
  Matches    5;  Conservative    5;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLH 13
              | |:| :::|:|
Db         44 GIGLAATQVNVH 55


Search completed: May 13, 2006, 08:14:40
Job time : 42 secs
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GenCore version 5.1.8

OM protein - protein search, using sw model

Run on: May 13, 2006, 08:08:53 ; Search time 226 Seconds
(without alignments)
43.705 Million cell updates/sec

Title: US-10-769-514-17
Perfect score: 74
Sequence: 1 MGYGMALSKINLHN 14

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2166443 seqs, 705528306 residues

Total number of hits satisfying chosen parameters: 2166443

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 100 summaries

Database : UniProt_05.80:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	74	100.0	702	2	O85050_MORCA	O85050 moraxella c
2	65	87.8	706	2	O85052_MORCA	O85052 moraxella c
3	63	85.1	714	2	O85056_MORCA	O85056 moraxella c
4	46	62.2	188	1	YAAH_ECOLI	P28695 escherichia
5	46	62.2	188	2	Q57TP6_SALCH	Q57tp6 salmonella
6	46	62.2	188	2	Q5PDN1_SALPA	Q5pdn1 salmonella
7	46	62.2	188	2	Q7CRA2_SALTY	Q7cra2 salmonella
8	46	62.2	188	2	Q8FLC8_ECOL6	Q8flc8 escherichia
9	46	62.2	188	2	Q8XGB2_SALTI	Q8xgb2 salmonella
10	46	62.2	189	2	Q7N8Y5_PHOLL	Q7n8y5 photorhabdu
11	46	62.2	191	2	Q6D0B6_ERWCT	Q6d0b6 erwinia car
12	46	62.2	196	2	Q74Q11_YERPE	Q74q11 yersinia pe
13	46	62.2	197	2	Q6M0C3_METMP	Q6m0c3 methanococc
14	46	62.2	203	2	Q8ZIM8_YERPE	Q8zim8 yersinia pe
15	46	62.2	203	2	Q66ET1_YERPS	Q66et1 yersinia ps
16	46	62.2	851	2	Q5CQ75_CRYPV	Q5cq75 cryptospori
17	44	59.5	183	2	Q727R6_DESVH	Q727r6 desulfovibr
18	44	59.5	185	2	Q6AK78_DESPS	Q6ak78 desulfotale
19	44	59.5	186	2	Q7P0P5_CHRVO	Q7p0p5 chromobacte
20	44	59.5	189	2	Q8EBD7_SHEON	Q8ebd7 shewanella
21	44	59.5	196	2	Q87S04_VIBPA	Q87s04 vibrio para
22	44	59.5	197	2	Q6LPH7_PHOPR	Q6lph7 photobacter
23	44	59.5	197	2	Q8DF09_VIBVU	Q8df09 vibrio vuln
24	44	59.5	197	2	Q9KTW0_VIBCH	Q9ktw0 vibrio chol
25	44	59.5	197	2	Q7MND9_VIBVY	Q7mnd9 vibrio vuln
26	44	59.5	200	2	Q8PYF9_METMA	Q8pyf9 methanosarc
27	44	59.5	201	2	Q8TIY1_METAC	Q8tiy1 methanosarc
28	44	59.5	343	2	Q4UYJ0_XANCP	Q4uyj0 xanthomonas
29	44	59.5	343	2	Q8P5I5_XANCP	Q8p5i5 xanthomonas
30	44	59.5	362	2	Q6MST2_MYCMS	Q6mst2 mycoplasma
31	44	59.5	501	2	Q4H6L8_9DEIO	Q4h6l8 deinococcus
32	44	59.5	694	2	Q4RYL2_TETNG	Q4ryl2 tetraodon n
33	44	59.5	887	2	Q899T2_CLOTE	Q899t2 clostridium
34	43	58.1	102	1	RS24_AERPE	Q9ycy0 aeropyrum p
35	43	58.1	214	2	Q4NQB2_9DELT	Q4nqb2 anaeromyxob
36	42	56.8	541	2	Q4IKZ0_GIBZE	Q4ikz0 gibberella
37	42	56.8	696	1	SLIK1_HUMAN	Q96px8 homo sapien
38	42	56.8	696	1	SLIK1_MOUSE	Q810c1 mus musculu
39	42	56.8	696	2	Q5U5I6_HUMAN	Q5u5i6 homo sapien
40	42	56.8	696	2	Q5RAC4_PONPY	Q5rac4 pongo pygma
41	42	56.8	760	2	Q69Z70_MOUSE	Q69z70 mus musculu
42	42	56.8	918	1	Y011_CLOPE	Q8xpf2 clostridium
43	41	55.4	303	2	Q9N6G9_LEIMA	Q9n6g9 leishmania
44	41	55.4	363	2	Q74EF8_GEOSL	Q74ef8 geobacter s
45	41	55.4	370	2	Q9N686_LEIMA	Q9n686 leishmania
46	41	55.4	478	2	Q8MXL0_LEIMA	Q8mxl0 leishmania
47	41	55.4	894	2	Q74YT9_YERPE	Q74yt9 yersinia pe
48	41	55.4	906	1	Y010_CLOAB	Q97n28 clostridium
49	41	55.4	1472	2	Q8A936_BACTN	Q8a936 bacteroides
50	40	54.1	195	2	Q5E5Q1_VIBF1	Q5e5q1 vibrio fisc
51	40	54.1	216	2	Q9FY32_DIGLA	Q9fy32 digitalis l
52	40	54.1	310	2	Q93L64_BACTI	Q93l64 bacillus th
53	40	54.1	423	2	Q58EH8_BRARE	Q58eh8 brachydanio
54	40	54.1	429	1	MTA1_ACEPA	O52702 acetobacter
55	40	54.1	429	2	P70750_ACEPA	P70750 acetobacter
56	40	54.1	432	1	HEM1_XYLFA	Q9pa72 xylella fas
57	40	54.1	432	1	HEM1_XYLFT	Q87a18 xylella fas
58	40	54.1	569	2	Q8KNX1_BACTI	Q8knx1 bacillus th
59	40	54.1	569	2	Q74NY4_BACC1	Q74ny4 bacillus ce
60	40	54.1	607	2	Q9K5R2_BACHD	Q9k5r2 bacillus ha
61	40	54.1	665	2	Q4RXK0_TETNG	Q4rxk0 tetraodon n
62	40	54.1	753	2	Q83AL7_COXBU	Q83al7 coxiella bu
63	40	54.1	1122	2	Q23682_CAEEL	Q23682 caenorhabdi
64	40	54.1	7214	2	Q5AUZ6_EMENI	Q5auz6 aspergillus
65	39.5	53.4	1854	2	Q63131_RAT	Q63131 rattus norv
66	39.5	53.4	2317	2	Q63130_RAT	Q63130 rattus norv
67	39.5	53.4	2338	2	Q63132_RAT	Q63132 rattus norv
68	39	52.7	89	1	USG_CAUCR	P12288 caulobacter
69	39	52.7	204	1	Y215_METTH	O26317 methanobact
70	39	52.7	343	2	Q8PGX5_XANAC	Q8pgx5 xanthomonas
71	39	52.7	368	2	Q72KY8_THET2	Q72ky8 thermus the
72	39	52.7	380	2	Q7RP13_PLAYO	Q7rp13 plasmodium
73	39	52.7	406	2	Q5H3W6_XANOR	Q5h3w6 xanthomonas
74	39	52.7	407	2	Q9AEU0_STRGN	Q9aeu0 streptococc
75	39	52.7	444	2	Q5LDQ1_BACFN	Q5ldq1 bacteroides
76	39	52.7	487	2	Q5M8Z9_XENTR	Q5m8z9 xenopus tro
77	39	52.7	521	2	Q5X8G4_LEGPA	Q5x8g4 legionella
78	39	52.7	521	2	Q5ZYY3_LEGPH	Q5zyy3 legionella
79	39	52.7	580	2	Q7QHE9_ANOGA	Q7qhe9 anopheles g
80	39	52.7	615	2	Q600M4_MYCHY	Q600m4 mycoplasma
81	39	52.7	919	2	Q4UBP6_THEAN	Q4ubp6 theileria a
82	39	52.7	1032	1	Y1060_SYNY3	P72637 synechocyst
83	39	52.7	6256	2	Q4U446_POLCB	Q4u446 polyangium
84	38.5	52.0	518	2	Q897Y2_CLOTE	Q897y2 clostridium
85	38	51.4	105	2	Q64EA9_9ARCH	Q64ea9 uncultured
86	38	51.4	132	2	Q6LYY7_METMP	Q6lyy7 methanococc
87	38	51.4	171	2	Q6SKZ5_9CRUS	Q6skz5 hutchinsoni
88	38	51.4	181	2	Q4K7U4_PSEF5	Q4k7u4 pseudomonas
89	38	51.4	194	2	Q5QV66_IDILO	Q5qv66 idiomarina
90	38	51.4	196	2	Q7SX89_BRARE	Q7sx89 brachydanio
91	38	51.4	206	2	Q4WBP5_ASPFU	Q4wbp5 aspergillus
92	38	51.4	251	2	Q883W8_PSESM	Q883w8 pseudomonas
93	38	51.4	273	2	Q9M501_9LILI	Q9m501 dioscorea a
94	38	51.4	279	2	Q6FRE4_CANGA	Q6fre4 candida gla
95	38	51.4	282	2	Q97FS1_CLOAB	Q97fs1 clostridium
96	38	51.4	290	2	Q6KYW8_PICTO	Q6kyw8 picrophilus
97	38	51.4	291	2	Q4RF21_TETNG	Q4rf21 tetraodon n
98	38	51.4	316	2	Q48999_MYCCA	Q48999 mycoplasma
99	38	51.4	366	2	Q4L6Q9_STAHJ	Q4l6q9 staphylococ
100	38	51.4	391	2	Q4UIL9_THEAN	Q4uil9 theileria a

ALIGNMENTS

```
RESULT 1
O85050_MORCA
ID   O85050_MORCA PRELIMINARY;      PRT;   702 AA.
AC   O85050;
DT   01-NOV-1998 (TrEMBLrel. 08, Created)
DT   01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Transferrin binding protein B.
GN   Name=tbpB;
OS   Moraxella catarrhalis.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
OC   Moraxellaceae; Moraxella.
OX   NCBI_TaxID=480;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=4223;
RX   MEDLINE=98380363; PubMed=9712766;
RA   Myers L.E., Yang Y.p., Du R.p., Wang Q., Harkness R.E.,
RA   Schryvers A.B., Klein M.H., Loosmore S.M.;
RT   "The transferrin binding protein B of Moraxella catarrhalis elicits
RT   bactericidal antibodies and is a potential vaccine antigen.";
RL   Infect. Immun. 66:4183-4192(1998).
DR   EMBL; AF039312; AAC34277.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   GO; GO:0004998; F:transferrin receptor activity; IEA.
DR   InterPro; IPR001677; Transferrin_bind.
DR   Pfam; PF01298; Lipoprotein_5; 1.
SQ   SEQUENCE   702 AA;  75458 MW;  974CBC14BBB0555A CRC64;

  Query Match             100.0%;  Score 74;  DB 2;  Length 702;
  Best Local Similarity   100.0%;  Pred. No. 0.00019;
  Matches   14;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||||
Db        100 MGYGMALSKINLHN 113


RESULT 2
O85052_MORCA
ID   O85052_MORCA PRELIMINARY;      PRT;   706 AA.
AC   O85052;
DT   01-NOV-1998 (TrEMBLrel. 08, Created)
DT   01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Transferrin binding protein B.
GN   Name=tbpB;
OS   Moraxella catarrhalis.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
OC   Moraxellaceae; Moraxella.
OX   NCBI_TaxID=480;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=M35;
RX   MEDLINE=98380363; PubMed=9712766;
RA   Myers L.E., Yang Y.p., Du R.p., Wang Q., Harkness R.E.,
RA   Schryvers A.B., Klein M.H., Loosmore S.M.;
RT   "The transferrin binding protein B of Moraxella catarrhalis elicits
RT   bactericidal antibodies and is a potential vaccine antigen.";
RL   Infect. Immun. 66:4183-4192(1998).
DR   EMBL; AF039314; AAC34279.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   GO; GO:0004998; F:transferrin receptor activity; IEA.
DR   InterPro; IPR001677; Transferrin_bind.
DR   Pfam; PF01298; Lipoprotein_5; 1.
SQ   SEQUENCE   706 AA;  76492 MW;  FB817A285958E64B CRC64;

  Query Match             87.8%;  Score 65;  DB 2;  Length 706;
  Best Local Similarity   92.3%;  Pred. No. 0.0082;
  Matches   12;  Conservative    1;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLH 13
              |||||||||:|||
Db        100 MGYGMALSKMNLH 112


RESULT 3
O85056_MORCA
ID   O85056_MORCA PRELIMINARY;      PRT;   714 AA.
AC   O85056;
DT   01-NOV-1998 (TrEMBLrel. 08, Created)
DT   01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Transferrin binding protein B.
GN   Name=tbpB;
OS   Moraxella catarrhalis.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
OC   Moraxellaceae; Moraxella.
OX   NCBI_TaxID=480;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=R1;
RX   MEDLINE=98380363; PubMed=9712766;
RA   Myers L.E., Yang Y.p., Du R.p., Wang Q., Harkness R.E.,
RA   Schryvers A.B., Klein M.H., Loosmore S.M.;
RT   "The transferrin binding protein B of Moraxella catarrhalis elicits
RT   bactericidal antibodies and is a potential vaccine antigen.";
RL   Infect. Immun. 66:4183-4192(1998).
DR   EMBL; AF039316; AAC34283.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   GO; GO:0004998; F:transferrin receptor activity; IEA.
DR   InterPro; IPR001677; Transferrin_bind.
DR   Pfam; PF01298; Lipoprotein_5; 1.
SQ   SEQUENCE   714 AA;  76846 MW;  F5B1174C4815E4EA CRC64;

  Query Match             85.1%;  Score 63;  DB 2;  Length 714;
  Best Local Similarity   85.7%;  Pred. No. 0.019;
  Matches   12;  Conservative    2;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||||||||||||::
Db        100 MGYGMALSKINLYD 113


RESULT 4
YAAH_ECOLI
ID   YAAH_ECOLI     STANDARD;      PRT;   188 AA.
AC   P28695; Q8KMY2;
DT   01-DEC-1992 (Rel. 24, Created)
DT   01-DEC-1992 (Rel. 24, Last sequence update)
DT   13-SEP-2005 (Rel. 48, Last annotation update)
DE   Inner membrane protein yaaH.
GN   Name=yaaH; OrderedLocusNames=b0010, z0010, ECs0010, SF0011, S0010;
OS   Escherichia coli,
OS   Escherichia coli O157:H7, and
OS   Shigella flexneri.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
OC   Enterobacteriaceae; Escherichia.
OX   NCBI_TaxID=562, 83334, 623;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   SPECIES=E.coli;
RX   MEDLINE=94003405; PubMed=8400364;
RA   James R., Dean D.O., Debbage J.;
RT   "Five open reading frames upstream of the dnaK gene of E. coli.";
RL   DNA Seq. 3:327-332(1993).
RN   [2]
RP   NUCLEOTIDE SEQUENCE.
RC   SPECIES=E.coli; STRAIN=K12;
RX   MEDLINE=92334977; PubMed=1630901;
RA   Yura T., Mori H., Nagai H., Nagata T., Ishihama A., Fujita N.,
RA   Isono K., Mizobuchi K., Nakata A.;
RT   "Systematic sequencing of the Escherichia coli genome: analysis of the
RT   0-2.4 min region.";
RL   Nucleic Acids Res. 20:3305-3308(1992).
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RN   [3]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC   SPECIES=E.coli; STRAIN=K12 / MG1655;
RX   MEDLINE=97426617; PubMed=9278503; DOI=10.1126/science.277.5331.1453;
RA   Blattner F.R., Plunkett G. III, Bloch C.A., Perna N.T., Burland V.,
RA   Riley M., Collado-Vides J., Glasner J.D., Rode C.K., Mayhew G.F.,
RA   Gregor J., Davis N.W., Kirkpatrick H.A., Goeden M.A., Rose D.J.,
RA   Mau B., Shao Y.;
RT   "The complete genome sequence of Escherichia coli K-12.";
RL   Science 277:1453-1474(1997).
RN   [4]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC   SPECIES=E.coli; STRAIN=O157:H7 / EDL933 / ATCC 700927 / EHEC;
RX   MEDLINE=21074935; PubMed=11206551; DOI=10.1038/35054089;
RA   Perna N.T., Plunkett G. III, Burland V., Mau B., Glasner J.D.,
RA   Rose D.J., Mayhew G.F., Evans P.S., Gregor J., Kirkpatrick H.A.,
RA   Posfai G., Hackett J., Klink S., Boutin A., Shao Y., Miller L.,
RA   Grotbeck E.J., Davis N.W., Lim A., Dimalanta E.T., Potamousis K.,
RA   Apodaca J., Anantharaman T.S., Lin J., Yen G., Schwartz D.C.,
RA   Welch R.A., Blattner F.R.;
RT   "Genome sequence of enterohaemorrhagic Escherichia coli O157:H7.";
RL   Nature 409:529-533(2001).
RN   [5]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC   SPECIES=E.coli; STRAIN=O157:H7 / Sakai / RIMD 0509952 / EHEC;
RX   MEDLINE=21156231; PubMed=11258796;
RA   Hayashi T., Makino K., Ohnishi M., Kurokawa K., Ishii K., Yokoyama K.,
RA   Han C.-G., Ohtsubo E., Nakayama K., Murata T., Tanaka M., Tobe T.,
RA   Iida T., Takami H., Honda T., Sasakawa C., Ogasawara N., Yasunaga T.,
RA   Kuhara S., Shiba T., Hattori M., Shinagawa H.;
RT   "Complete genome sequence of enterohemorrhagic Escherichia coli
RT   O157:H7 and genomic comparison with a laboratory strain K-12.";
RL   DNA Res. 8:11-22(2001).
RN   [6]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC   SPECIES=S.flexneri; STRAIN=301 / Serotype 2a;
RX   MEDLINE=22272406; PubMed=12384590; DOI=10.1093/nar/gkf566;
RA   Jin Q., Yuan Z., Xu J., Wang Y., Shen Y., Lu W., Wang J., Liu H.,
RA   Yang J., Yang F., Zhang X., Zhang J., Yang G., Wu H., Qu D., Dong J.,
RA   Sun L., Xue Y., Zhao A., Gao Y., Zhu J., Kan B., Ding K., Chen S.,
RA   Cheng H., Yao Z., He B., Chen R., Ma D., Qiang B., Wen Y., Hou Y.,
RA   Yu J.;
RT   "Genome sequence of Shigella flexneri 2a: insights into pathogenicity
RT   through comparison with genomes of Escherichia coli K12 and O157.";
RL   Nucleic Acids Res. 30:4432-4441(2002).
RN   [7]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC   SPECIES=S.flexneri; STRAIN=2457T / ATCC 700930 / Serotype 2a;
RX   MEDLINE=22590274; PubMed=12704152;
RX   DOI=10.1128/IAI.71.5.2775-2786.2003;
RA   Wei J., Goldberg M.B., Burland V., Venkatesan M.M., Deng W.,
RA   Fournier G., Mayhew G.F., Plunkett G. III, Rose D.J., Darling A.,
RA   Mau B., Perna N.T., Payne S.M., Runyen-Janecky L.J., Zhou S.,
RA   Schwartz D.C., Blattner F.R.;
RT   "Complete genome sequence and comparative genomics of Shigella
RT   flexneri serotype 2a strain 2457T.";
RL   Infect. Immun. 71:2775-2786(2003).
RN   [8]
RP   TOPOLOGY.
RC   SPECIES=E.coli; STRAIN=K12 / MG1655;
RX   PubMed=15919996; DOI=10.1126/science.1109730;
RA   Daley D.O., Rapp M., Granseth E., Melen K., Drew D., von Heijne G.;
RT   "Global topology analysis of the Escherichia coli inner membrane
RT   proteome.";
RL   Science 308:1321-1323(2005).
CC   -!- SUBCELLULAR LOCATION: Integral membrane protein. Inner membrane.
CC   -!- SIMILARITY: Belongs to the GPR1/FUN34/yaaH family.
CC   ----------------------------------------------------------------------
CC   This Swiss-Prot entry is copyright. It is produced through a collaboration
CC   between the Swiss Institute of Bioinformatics and the EMBL outstation -
CC   the European Bioinformatics Institute. There are no restrictions on its
CC   use as long as its content is in no way modified and this statement is not
CC   removed.
CC   ----------------------------------------------------------------------
DR   EMBL; X67700; CAA47931.1; -; Genomic_DNA.
DR   EMBL; D10483; BAB96588.1; -; Genomic_DNA.
DR   EMBL; U00096; AAC73121.1; -; Genomic_DNA.
DR   EMBL; AE005174; AAG54310.1; -; Genomic_DNA.
DR   EMBL; BA000007; BAB33433.1; -; Genomic_DNA.
DR   EMBL; AE005674; AAN41677.1; -; Genomic_DNA.
DR   EMBL; AE016978; AAP15556.1; -; Genomic_DNA.
DR   PIR; B85481; B85481.
DR   PIR; B90630; B90630.
DR   PIR; E56688; E56688.
DR   EchoBASE; EB1474; -.
DR   EcoGene; EG11512; yaaH.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome; Inner membrane; Membrane; Transmembrane.
FT   TOPO_DOM      1     13       Cytoplasmic (Potential).
FT   TRANSMEM     14     34       Potential.
FT   TOPO_DOM     35     35       Periplasmic (Potential).
FT   TRANSMEM     36     56       Potential.
FT   TOPO_DOM     57     63       Cytoplasmic (Potential).
FT   TRANSMEM     64     84       Potential.
FT   TOPO_DOM     85     97       Periplasmic (Potential).
FT   TRANSMEM     98    118       Potential.
FT   TOPO_DOM    119    122       Cytoplasmic (Potential).
FT   TRANSMEM    123    143       Potential.
FT   TOPO_DOM    144    148       Periplasmic (Potential).
FT   TRANSMEM    149    169       Potential.
FT   TOPO_DOM    170    188       Cytoplasmic (Potential).
FT   CONFLICT    111    111       L -> V (in Ref. 2).
SQ   SEQUENCE   188 AA;  20071 MW;  972101DD5949EBF4 CRC64;

  Query Match             62.2%;  Score 46;  DB 1;  Length 188;
  Best Local Similarity   57.1%;  Pred. No. 6.4;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTILLNLHN 28


RESULT 5
Q57TP6_SALCH
ID   Q57TP6_SALCH PRELIMINARY;      PRT;   188 AA.
AC   Q57TP6;
DT   10-MAY-2005 (TrEMBLrel. 30, Created)
DT   10-MAY-2005 (TrEMBLrel. 30, Last sequence update)
DT   10-MAY-2005 (TrEMBLrel. 30, Last annotation update)
DE   Putative regulator.
GN   Name=yaaH; OrderedLocusNames=SC0009;
OS   Salmonella cholerae-suis (Salmonella enterica).
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
OC   Enterobacteriaceae; Salmonella.
OX   NCBI_TaxID=591;
RN   [1]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC   STRAIN=SC-B67;
RX   PubMed=15781495;
RA   Chiu C.-H., Tang P., Chu C., Hu S., Bao Q., Yu J., Chou Y.-Y.,
RA   Wang H.-S., Lee Y.-S.;
RT   "The genome sequence of Salmonella enterica serovar Choleraesuis, a
RT   highly invasive and resistant zoonotic pathogen.";
RL   Nucleic Acids Res. 33:1690-1698(2005).
DR   EMBL; AE017220; AAX63915.1; -; Genomic_DNA.
KW   Complete proteome.
SQ   SEQUENCE   188 AA;  19933 MW;  4D655AC780BBB808 CRC64;

  Query Match             62.2%;  Score 46;  DB 2;  Length 188;
  Best Local Similarity   57.1%;  Pred. No. 6.4;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;
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Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTILLNLHN 28


RESULT 6
Q5PDN1_SALPA
ID   Q5PDN1_SALPA PRELIMINARY;      PRT;   188 AA.
AC   Q5PDN1;
DT   01-FEB-2005 (TrEMBLrel. 29, Created)
DT   01-FEB-2005 (TrEMBLrel. 29, Last sequence update)
DT   01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
DE   Hypothetical protein yaaH.
GN   Name=yaaH; OrderedLocusNames=SPA0009;
OS   Salmonella paratyphi-a.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
OC   Enterobacteriaceae; Salmonella.
OX   NCBI_TaxID=54388;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=ATCC 9150;
RX   PubMed=15531882; DOI=10.1038/ng1470;
RA   McClelland M., Sanderson K.E., Clifton S.W., Latreille P.,
RA   Porwollik S., Sabo A., Meyer R., Bieri T., Ozersky P., McLellan M.,
RA   Harkins C.R., Wang C., Nguyen C., Berghoff A., Elliott G.,
RA   Kohlberg S., Strong C., Du F., Carter J., Kremizki C., Layman D.,
RA   Leonard S., Sun H., Fulton L., Nash W., Miner T., Minx P.,
RA   Delehaunty K., Fronick C., Magrini V., Nhan M., Warren W., Florea L.,
RA   Spieth J., Wilson R.K.;
RT   "Comparison of genome degradation in Paratyphi A and Typhi, human-
RT   restricted serovars of Salmonella enterica that cause typhoid.";
RL   Nat. Genet. 36:1268-1274(2004).
DR   EMBL; CP000026; AAV76047.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome; Hypothetical protein.
SQ   SEQUENCE   188 AA;  19962 MW;  9CE9B7D3D1EEAD41 CRC64;

  Query Match           62.2%;  Score 46;  DB 2;  Length 188;
  Best Local Similarity   57.1%;  Pred. No. 6.4;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTILLNLHN 28


RESULT 7
Q7CRA2_SALTY
ID   Q7CRA2_SALTY PRELIMINARY;      PRT;   188 AA.
AC   Q7CRA2;
DT   05-JUL-2004 (TrEMBLrel. 27, Created)
DT   05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT   05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE   Putative regulatory protein.
GN   Name=yaaH; OrderedLocusNames=STM0009;
OS   Salmonella typhimurium.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
OC   Enterobacteriaceae; Salmonella.
OX   NCBI_TaxID=602;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=LT2;
RX   MEDLINE=21534948; PubMed=11677609; DOI=10.1038/35101614;
RA   McClelland M., Sanderson K.E., Spieth J., Clifton S.W., Latreille P.,
RA   Courtney L., Porwollik S., Ali J., Dante M., Du F., Hou S., Layman D.,
RA   Leonard S., Nguyen C., Scott K., Holmes A., Grewal N., Mulvaney E.,
RA   Ryan E., Sun H., Florea L., Miller W., Stoneking T., Nhan M.,
RA   Waterston R., Wilson R.K.;
RT   "Complete genome sequence of Salmonella enterica serovar Typhimurium
RT   LT2.";
RL   Nature 413:852-856(2001).
DR   EMBL; AE008693; AAL18973.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome.
SQ   SEQUENCE   188 AA;  19933 MW;  4D655AC780BBB808 CRC64;

  Query Match           62.2%;  Score 46;  DB 2;  Length 188;
  Best Local Similarity   57.1%;  Pred. No. 6.4;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTILLNLHN 28


RESULT 8
Q8FLC8_ECOL6
ID   Q8FLC8_ECOL6 PRELIMINARY;      PRT;   188 AA.
AC   Q8FLC8;
DT   01-MAR-2003 (TrEMBLrel. 23, Created)
DT   01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
DT   01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE   Hypothetical protein yaaH.
GN   Name=yaaH; OrderedLocusNames=c0015;
OS   Escherichia coli O6.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
OC   Enterobacteriaceae; Escherichia.
OX   NCBI_TaxID=217992;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=O6:H1 / CFT073 / ATCC 700928;
RX   MEDLINE=22388234; PubMed=12471157; DOI=10.1073/pnas.252529799;
RA   Welch R.A., Burland V., Plunkett G. III, Redford P., Roesch P.,
RA   Rasko D., Buckles E.L., Liou S.-R., Boutin A., Hackett J., Stroud D.,
RA   Mayhew G.F., Rose D.J., Zhou S., Schwartz D.C., Perna N.T.,
RA   Mobley H.L.T., Donnenberg M.S., Blattner F.R.;
RT   "Extensive mosaic structure revealed by the complete genome sequence
RT   of uropathogenic Escherichia coli.";
RL   Proc. Natl. Acad. Sci. U.S.A. 99:17020-17024(2002).
DR   EMBL; AE016755; AAN78515.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome; Hypothetical protein.
SQ   SEQUENCE   188 AA;  20085 MW;  815BBD1B23F53BF4 CRC64;

  Query Match           62.2%;  Score 46;  DB 2;  Length 188;
  Best Local Similarity   57.1%;  Pred. No. 6.4;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTILLNLHN 28


RESULT 9
Q8XGB2_SALTI
ID   Q8XGB2_SALTI PRELIMINARY;      PRT;   188 AA.
AC   Q8XGB2; Q7ANM8;
DT   01-MAR-2002 (TrEMBLrel. 20, Created)
DT   01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
DT   13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
DE   Hypothetical protein yaaH (Hypothetical protein STY0009).
GN   Name=yaaH; OrderedLocusNames=STY0009, t0009;
OS   Salmonella typhi.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
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OC   Enterobacteriaceae; Salmonella.
OX   NCBI_TaxID=601;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=Ty2 / ATCC 700931;
RX   MEDLINE=22531367; PubMed=12644504;
RX   DOI=10.1128/JB.185.7.2330-2337.2003;
RA   Deng W., Liou S.-R., Plunkett G. III, Mayhew G.F., Rose D.J.,
RA   Burland V., Kodoyianni V., Schwartz D.C., Blattner F.R.;
RT   "Comparative genomics of Salmonella enterica serovar Typhi strains Ty2
RT   and CT18.";
RL   J. Bacteriol. 185:2330-2337(2003).
RN   [2]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=CT18;
RX   MEDLINE=21534947; PubMed=11677608; DOI=10.1038/35101607;
RA   Parkhill J., Dougan G., James K.D., Thomson N.R., Pickard D., Wain J.,
RA   Churcher C.M., Mungall K.L., Bentley S.D., Holden M.T.G., Sebaihia M.,
RA   Baker S., Basham D., Brooks K., Chillingworth T., Connerton P.,
RA   Cronin A., Davis P., Davies R.M., Dowd L., White N., Farrar J.,
RA   Feltwell T., Hamlin N., Haque A., Hien T.T., Holroyd S., Jagels K.,
RA   Krogh A., Larsen T.S., Leather S., Moule S., O'Gaora P., Parry C.,
RA   Quail M.A., Rutherford K.M., Simmonds M., Skelton J., Stevens K.,
RA   Whitehead S., Barrell B.G.;
RT   "Complete genome sequence of a multiple drug resistant Salmonella
RT   enterica serovar Typhi CT18.";
RL   Nature 413:848-852(2001).
DR   EMBL; AE016834; AAO67743.1; -; Genomic_DNA.
DR   EMBL; AL627265; CAD01162.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome; Hypothetical protein.
SQ   SEQUENCE   188 AA;  19933 MW;  4D655AC780BBB808 CRC64;

  Query Match             62.2%;  Score 46;  DB 2;  Length 188;
  Best Local Similarity   57.1%;  Pred. No. 6.4;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTILLNLHN 28


RESULT 10
Q7N8Y5_PHOLL
ID   Q7N8Y5_PHOLL PRELIMINARY;      PRT;   189 AA.
AC   Q7N8Y5;
DT   01-MAR-2004 (TrEMBLrel. 26, Created)
DT   01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Similar to unknown protein YaaH of Escherichia coli.
GN   OrderedLocusNames=plu0578;
OS   Photorhabdus luminescens (subsp. laumondii).
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
OC   Enterobacteriaceae; Photorhabdus.
OX   NCBI_TaxID=141679;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=TT01;
RX   MEDLINE=22957627; PubMed=14528314; DOI=10.1038/nbt886;
RA   Duchaud E., Rusniok C., Frangeul L., Buchrieser C., Givaudan A.,
RA   Taourit S., Bocs S., Boursaux-Eude C., Chandler M., Charles J.-F.,
RA   Dassa E., Derose R., Derzelle S., Freyssinet G., Gaudriault S.,
RA   Medigue C., Lanois A., Powell K., Siguier P., Vincent R., Wingate V.,
RA   Zouine M., Glaser P., Boemare N., Danchin A., Kunst F.;
RT   "The genome sequence of the entomopathogenic bacterium Photorhabdus
RT   luminescens.";
RL   Nat. Biotechnol. 21:1307-1313(2003).
DR   EMBL; BX571860; CAE12873.1; -; Genomic_DNA.
DR   PhotoList; plu0578; -.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
KW   Complete proteome.
SQ   SEQUENCE   189 AA;  20289 MW;  E9D5C44D35306A7F CRC64;

  Query Match             62.2%;  Score 46;  DB 2;  Length 189;
  Best Local Similarity   57.1%;  Pred. No. 6.4;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTILLNLHN 28


RESULT 11
Q6D0B6_ERWCT
ID   Q6D0B6_ERWCT PRELIMINARY;      PRT;   191 AA.
AC   Q6D0B6;
DT   25-OCT-2004 (TrEMBLrel. 28, Created)
DT   25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
DT   25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE   Putative membrane protein.
GN   OrderedLocusNames=ECA3883;
OS   Erwinia carotovora (subsp. atroseptica) (Pectobacterium atrosepticum).
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
OC   Enterobacteriaceae; Pectobacterium.
OX   NCBI_TaxID=29471;
RN   [1]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC   STRAIN=SCRI 1043 / ATCC BAA-672;
RX   PubMed=15263089; DOI=10.1073/pnas.0402424101;
RA   Bell K.S., Sebaihia M., Pritchard L., Holden M.T.G., Hyman L.J.,
RA   Holeva M.C., Thomson N.R., Bentley S.D., Churcher L.J.C., Mungall K.,
RA   Atkin R., Bason N., Brooks K., Chillingworth T., Clark K., Doggett J.,
RA   Fraser A., Hance Z., Hauser H., Jagels K., Moule S., Norbertczak H.,
RA   Ormond D., Price C., Quail M.A., Sanders M., Walker D., Whitehead S.,
RA   Salmond G.P.C., Birch P.R.J., Parkhill J., Toth I.K.;
RT   "Genome sequence of the enterobacterial phytopathogen Erwinia
RT   carotovora subsp. atroseptica and characterization of virulence
RT   factors.";
RL   Proc. Natl. Acad. Sci. U.S.A. 101:11105-11110(2004).
DR   EMBL; BX950851; CAG76781.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
KW   Complete proteome.
SQ   SEQUENCE   191 AA;  20218 MW;  475605990E163AC1 CRC64;

  Query Match             62.2%;  Score 46;  DB 2;  Length 191;
  Best Local Similarity   57.1%;  Pred. No. 6.5;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTILLNLHN 28


RESULT 12
Q74Q11_YERPE
ID   Q74Q11_YERPE PRELIMINARY;      PRT;   196 AA.
AC   Q74Q11;
DT   05-JUL-2004 (TrEMBLrel. 27, Created)
DT   05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT   05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE   Putative membrane protein.
GN   OrderedLocusNames=YP3713;
OS   Yersinia pestis.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
OC   Enterobacteriaceae; Yersinia.
OX   NCBI_TaxID=632;
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RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=91001;
RX   PubMed=15368893;
RA   Song Y., Tong Z., Wang J., Wang L., Guo Z., Han Y., Zhang J., Pei D.,
RA   Zhou D., Qin H., Pang X., Han Y., Zhai J., Li M., Cui B., Qi Z.,
RA   Jin L., Dai R., Chen F., Li S., Ye C., Du Z., Lin W., Wang J., Yu J.,
RA   Yang H., Wang J., Huang P., Yang R.;
RT   "Complete genome sequence of Yersinia pestis strain 91001, an isolate
RT   avirulent to humans.";
RL   DNA Res. 11:179-197(2004).
DR   EMBL; AE017141; AAS63861.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
SQ   SEQUENCE   196 AA;  21046 MW;  B74D3F30EDC2047D CRC64;

  Query Match             62.2%;  Score 46;  DB 2;  Length 196;
  Best Local Similarity   57.1%;  Pred. No. 6.6;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTVLLNLHN 28


RESULT 13
Q6M0C3_METMP
ID   Q6M0C3_METMP PRELIMINARY;      PRT;   197 AA.
AC   Q6M0C3;
DT   05-JUL-2004 (TrEMBLrel. 27, Created)
DT   05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT   05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE   Hypothetical protein.
GN   OrderedLocusNames=MMP0348;
OS   Methanococcus maripaludis.
OC   Archaea; Euryarchaeota; Methanococci; Methanococcales;
OC   Methanococcaceae; Methanococcus.
OX   NCBI_TaxID=39152;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=S2 / LL;
RX   PubMed=15466049; DOI=10.1128/JB.186.20.6956-6969.2004;
RA   Hendrickson E.L., Kaul R., Zhou Y., Bovee D., Chapman P., Chung J.,
RA   Conway de Macario E., Dodsworth J.A., Gillett W., Graham D.E.,
RA   Hackett M., Haydock A.K., Kang A., Land M.L., Levy R., Lie T.J.,
RA   Major T.A., Moore B.C., Porat I., Palmeiri A., Rouse G.,
RA   Saenphimmachak C., Soell D., Van Dien S., Wang T., Whitman W.B.,
RA   Xia Q., Zhang Y., Larimer F.W., Olson M.V., Leigh J.A.;
RT   "Complete genome sequence of the genetically tractable
RT   hydrogenotrophic methanogen Methanococcus maripaludis.";
RL   J. Bacteriol. 186:6956-6969(2004).
DR   EMBL; BX957219; CAF29904.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome; Hypothetical protein.
SQ   SEQUENCE   197 AA;  21273 MW;  1C27A47FBB95A1BE CRC64;

  Query Match             62.2%;  Score 46;  DB 2;  Length 197;
  Best Local Similarity   57.1%;  Pred. No. 6.7;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         21 MGFGMTTVLLNLHN 34


RESULT 14
Q8ZIM8_YERPE
ID   Q8ZIM8_YERPE PRELIMINARY;      PRT;   203 AA.
AC   Q8ZIM8; Q7CG75;
DT   01-MAR-2002 (TrEMBLrel. 20, Created)
DT   01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
DT   01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
DE   Putative membrane protein (Hypothetical protein y3707).
GN   Name=yaaH; OrderedLocusNames=YPO0467, y3707;
OS   Yersinia pestis.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
OC   Enterobacteriaceae; Yersinia.
OX   NCBI_TaxID=632;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=CO-92 / Biovar Orientalis;
RX   MEDLINE=21470413; PubMed=11586360; DOI=10.1038/35097083;
RA   Parkhill J., Wren B.W., Thomson N.R., Titball R.W., Holden M.T.G.,
RA   Prentice M.B., Sebaihia M., James K.D., Churcher C.M., Mungall K.L.,
RA   Baker S., Basham D., Bentley S.D., Brooks K., Cerdeno-Tarraga A.-M.,
RA   Chillingworth T., Cronin A., Davies R.M., Davis P., Dougan G.,
RA   Feltwell T., Hamlin N., Holroyd S., Jagels K., Karlyshev A.V.,
RA   Leather S., Moule S., Oyston P.C.F., Quail M.A., Rutherford K.M.,
RA   Simmonds M., Skelton J., Stevens K., Whitehead S., Barrell B.G.;
RT   "Genome sequence of Yersinia pestis, the causative agent of plague.";
RL   Nature 413:523-527(2001).
RN   [2]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=KIM5 / Biovar Mediaevalis;
RX   MEDLINE=22137863; PubMed=12142430;
RX   DOI=10.1128/JB.184.16.4601-4611.2002;
RA   Deng W., Burland V., Plunkett G. III, Boutin A., Mayhew G.F., Liss P.,
RA   Perna N.T., Rose D.J., Mau B., Zhou S., Schwartz D.C.,
RA   Fetherston J.D., Lindler L.E., Brubaker R.R., Plano G.V.,
RA   Straley S.C., McDonough K.A., Nilles M.L., Matson J.S., Blattner F.R.,
RA   Perry R.D.;
RT   "Genome sequence of Yersinia pestis KIM.";
RL   J. Bacteriol. 184:4601-4611(2002).
DR   EMBL; AJ414142; CAC89323.1; -; Genomic_DNA.
DR   EMBL; AE013974; AAM87255.1; -; Genomic_DNA.
DR   PIR; AH0057; AH0057.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
KW   Complete proteome; Hypothetical protein.
SQ   SEQUENCE   203 AA;  21661 MW;  200282B6691CD661 CRC64;

  Query Match             62.2%;  Score 46;  DB 2;  Length 203;
  Best Local Similarity   57.1%;  Pred. No. 6.9;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTVLLNLHN 28


RESULT 15
Q66ET1_YERPS
ID   Q66ET1_YERPS PRELIMINARY;      PRT;   203 AA.
AC   Q66ET1;
DT   25-OCT-2004 (TrEMBLrel. 28, Created)
DT   25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
DT   25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE   Putative regulator; integral membrane protein.
GN   Name=yaaH; OrderedLocusNames=YPTB0610;
OS   Yersinia pseudotuberculosis.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
OC   Enterobacteriaceae; Yersinia.
OX   NCBI_TaxID=633;
RN   [1]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC   STRAIN=IP32953 / Serotype I;
RX   PubMed=15358858; DOI=10.1073/pnas.0404012101;
RA   Chain P.S.G., Carniel E., Larimer F.W., Lamerdin J., Stoutland P.O.,
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RA   Regala W.M., Georgescu A.M., Vergez L.M., Land M.L., Motin V.L.,
RA   Brubaker R.R., Fowler J., Hinnebusch J., Marceau M., Medigue C.,
RA   Simonet M., Chenal-Francisque V., Souza B., Dacheux D., Elliott J.M.,
RA   Derbise A., Hauser L.J., Garcia E.;
RT   "Insights into the evolution of Yersinia pestis through whole-genome
RT   comparison with Yersinia pseudotuberculosis.";
RL   Proc. Natl. Acad. Sci. U.S.A. 101:13826-13831(2004).
DR   EMBL; BX936398; CAH19850.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
KW   Complete proteome.
SQ   SEQUENCE   203 AA;  21644 MW;  A21F2175B2ABCA0A CRC64;

  Query Match             62.2%;  Score 46;  DB 2;  Length 203;
  Best Local Similarity   57.1%;  Pred. No. 6.9;
  Matches    8;  Conservative    2;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :||||
Db         15 MGFGMTTVLLNLHN 28


RESULT 16
Q5CQ75_CRYPV
ID   Q5CQ75_CRYPV PRELIMINARY;      PRT;   851 AA.
AC   Q5CQ75;
DT   10-MAY-2005 (TrEMBLrel. 30, Created)
DT   10-MAY-2005 (TrEMBLrel. 30, Last sequence update)
DT   10-MAY-2005 (TrEMBLrel. 30, Last annotation update)
DE   YprA. Lhr1/Ski2 family RNA SFII helicase (Fragment).
GN   ORFNames=cgd5_3870;
OS   Cryptosporidium parvum.
OC   Eukaryota; Alveolata; Apicomplexa; Coccidia; Eimeriida;
OC   Cryptosporidiidae; Cryptosporidium.
OX   NCBI_TaxID=5807;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=Iowa type II;
RX   PubMed=15044751; DOI=10.1126/science.1094786;
RA   Abrahamsen M.S., Templeton T.J., Enomoto S., Abrahante J.E., Zhu G.,
RA   Lancto C.A., Deng M., Liu C., Widmer G., Tzipori S., Buck G.A., Xu P.,
RA   Bankier A.T., Dear P.H., Konfortov B.A., Spriggs H.F., Iyer L.,
RA   Anantharaman V., Aravind L., Kapur V.;
RT   "Complete genome sequence of the apicomplexan, Cryptosporidium
RT   parvum.";
RL   Science 304:441-445(2004).
DR   EMBL; AAEE01000010; EAK87599.1; -; Genomic_DNA.
DR   GO; GO:0005524; F:ATP binding; IEA.
DR   GO; GO:0008026; F:ATP-dependent helicase activity; IEA.
DR   GO; GO:0016787; F:hydrolase activity; IEA.
DR   GO; GO:0003676; F:nucleic acid binding; IEA.
DR   InterPro; IPR001410; DEAD.
DR   InterPro; IPR011545; DEAD/DEAH_N.
DR   InterPro; IPR001650; Helicase_C.
DR   Pfam; PF00270; DEAD; 1.
DR   Pfam; PF00271; Helicase_C; 1.
DR   SMART; SM00487; DEXDc; 1.
DR   SMART; SM00490; HELICc; 1.
KW   ATP-binding; Helicase; Hydrolase.
FT   NON_TER       1       1
SQ   SEQUENCE   851 AA;  98508 MW;  1891EC62ADCF9D28 CRC64;

  Query Match             62.2%;  Score 46;  DB 2;  Length 851;
  Best Local Similarity   90.0%;  Pred. No. 28;
  Matches    9;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          5 MALSKINLHN 14
              ||| ||||||
Db        546 MALKKINLHN 555


RESULT 17
Q727R6_DESVH
ID   Q727R6_DESVH PRELIMINARY;      PRT;   183 AA.
AC   Q727R6;
DT   05-JUL-2004 (TrEMBLrel. 27, Created)
DT   05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT   05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE   Gpr1/Fun34/YaaH family protein.
GN   OrderedLocusNames=DVU2789;
OS   Desulfovibrio vulgaris (strain Hildenborough / ATCC 29579 / NCIMB
OS   8303).
OC   Bacteria; Proteobacteria; Deltaproteobacteria; Desulfovibrionales;
OC   Desulfovibrionaceae; Desulfovibrio.
OX   NCBI_TaxID=882;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RX   PubMed=15077118; DOI=10.1038/nbt959;
RA   Heidelberg J.F., Seshadri R., Haveman S.A., Hemme C.L., Paulsen I.T.,
RA   Kolonay J.F., Eisen J.A., Ward N.L., Methe B.A., Brinkac L.M.,
RA   Daugherty S.C., DeBoy R.T., Dodson R.J., Durkin A.S., Madupu R.,
RA   Nelson W.C., Sullivan S.A., Fouts D.E., Haft D.H., Selengut J.,
RA   Peterson J.D., Davidsen T.M., Zafar N., Zhou L., Radune D.,
RA   Dimitrov G., Hance M., Tran K., Khouri H.M., Gill J., Utterback T.R.,
RA   Feldblyum T.V., Wall J.D., Voordouw G., Fraser C.M.;
RT   "The genome sequence of the anaerobic, sulfate-reducing bacterium
RT   Desulfovibrio vulgaris Hildenborough.";
RL   Nat. Biotechnol. 22:554-559(2004).
DR   EMBL; AE017318; AAS97261.1; -; Genomic_DNA.
DR   TIGR; DVU2789; -.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome.
SQ   SEQUENCE   183 AA;  20134 MW;  82A9BDCDD18C4AA6 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 183;
  Best Local Similarity   50.0%;  Pred. No. 14;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         14 MGFGMTTVLLNIHN 27


RESULT 18
Q6AK78_DESPS
ID   Q6AK78_DESPS PRELIMINARY;      PRT;   185 AA.
AC   Q6AK78;
DT   25-OCT-2004 (TrEMBLrel. 28, Created)
DT   25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
DT   25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE   Conserved hypothetical membrane protein.
GN   OrderedLocusNames=DP2519;
OS   Desulfotalea psychrophila.
OC   Bacteria; Proteobacteria; Deltaproteobacteria; Desulfobacterales;
OC   Desulfobulbaceae; Desulfotalea.
OX   NCBI_TaxID=84980;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=LSv54 / DSM 12343;
RX   PubMed=15305914; DOI=10.1111/j.1462-2920.2004.00665.x;
RA   Rabus R., Ruepp A., Frickey T., Rattei T., Fartmann B., Stark M.,
RA   Bauer M., Zibat A., Lombardot T., Becker I., Amann J., Gellner K.,
RA   Teeling H., Leuschner W.D., Gloeckner F.-O., Lupas A.N., Amann R.,
RA   Klenk H.-P.;
RT   "The genome of Desulfotalea psychrophila, a sulfate-reducing bacterium
RT   from permanently cold Arctic sediments.";
RL   Environ. Microbiol. 6:887-902(2004).
DR   EMBL; CR522870; CAG37248.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
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DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
KW   Complete proteome.
SQ   SEQUENCE   185 AA;  20227 MW;  65AF87115A3B0560 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 185;
  Best Local Similarity   50.0%;  Pred. No. 14;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         15 MGFGMTTVLLNIHN 28


RESULT 19
Q7P0P5_CHRVO
ID   Q7P0P5_CHRVO PRELIMINARY;      PRT;   186 AA.
AC   Q7P0P5;
DT   01-MAR-2004 (TrEMBLrel. 26, Created)
DT   01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Probable membrane protein.
GN   OrderedLocusNames=CV0521;
OS   Chromobacterium violaceum.
OC   Bacteria; Proteobacteria; Betaproteobacteria; Neisseriales;
OC   Neisseriaceae; Chromobacterium.
OX   NCBI_TaxID=536;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=ATCC 12472 / DSM 30191;
RX   MEDLINE=22882880; PubMed=14500782; DOI=10.1073/pnas.1832124100;
RA   Vasconcelos A.T.R., de Almeida D.F., Hungria M., Guimaraes C.T.,
RA   Antonio R.V., Almeida F.C., de Almeida L.G.P., de Almeida R.,
RA   Alves-Gomes J.A., Andrade E.M., Araripe J., de Araujo M.F.F.,
RA   Astolfi-Filho S., Azevedo V., Baptista A.J., Bataus L.A.M.,
RA   Batista J.S., Belo A., van den Berg C., Bogo M., Bonatto S.,
RA   Bordignon J., Brigido M.M., Brito C.A., Brocchi M., Burity H.A.,
RA   Camargo A.A., Cardoso D.D.P., Carneiro N.P., Carraro D.M.,
RA   Carvalho C.M.B., Cascardo J.C.M., Cavada B.S., Chueire L.M.O.,
RA   Creczynski-Pasa T.B., Cunha-Junior N.C., Fagundes N., Falcao C.L.,
RA   Fantinatti F., Farias I.P., Felipe M.S.S., Ferrari L.P., Ferro J.A.,
RA   Ferro M.I.T., Franco G.R., Freitas N.S.A., Furlan L.R.,
RA   Gazzinelli R.T., Gomes E.A., Goncalves P.R., Grangeiro T.B.,
RA   Grattapaglia D., Grisard E.C., Hanna E.S., Jardim S.N., Laurino J.,
RA   Leoi L.C.T., Lima L.F.A., Loureiro M.F., Lyra M.C.C.P.,
RA   Madeira H.M.F., Manfio G.P., Maranhao A.Q., Martins W.S.,
RA   di Mauro S.M.Z., de Medeiros S.R.B., Meissner R.V., Moreira M.A.M.,
RA   Nascimento F.F., Nicolas M.F., Oliveira J.G., Oliveira S.C.,
RA   Paixao R.F.C., Parente J.A., Pedrosa F.O., Pena S.D.J., Pereira J.O.,
RA   Pereira M., Pinto L.S.R.C., Pinto L.S., Porto J.I.R., Potrich D.P.,
RA   Ramalho-Neto C.E., Reis A.M.M., Rigo L.U., Rondinelli E.,
RA   Santos E.B.P., Santos F.R., Schneider M.P.C., Seuanez H.N.,
RA   Silva A.M.R., da Silva A.L.C., Silva D.W., Silva R., Simoes I.C.,
RA   Simon D., Soares C.M.A., Soares R.B.A., Souza E.M., Souza K.R.L.,
RA   Souza R.C., Steffens M.B.R., Steindel M., Teixeira S.R., Urmenyi T.,
RA   Vettore A., Wassem R., Zaha A., Simpson A.J.G.;
RT   "The complete genome sequence of Chromobacterium violaceum reveals
RT   remarkable and exploitable bacterial adaptability.";
RL   Proc. Natl. Acad. Sci. U.S.A. 100:11660-11665(2003).
DR   EMBL; AE016911; AAQ58198.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome.
SQ   SEQUENCE   186 AA;  20086 MW;  177AB5FADFB92D56 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 186;
  Best Local Similarity   50.0%;  Pred. No. 15;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         14 MGFGMTTILLNIHN 27


RESULT 20
Q8EBD7_SHEON
ID   Q8EBD7_SHEON PRELIMINARY;      PRT;   189 AA.
AC   Q8EBD7;
DT   01-MAR-2003 (TrEMBLrel. 23, Created)
DT   01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
DT   01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE   Gpr1/fun34/yaaH family protein.
GN   OrderedLocusNames=SO3588;
OS   Shewanella oneidensis.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Alteromonadales;
OC   Shewanellaceae; Shewanella.
OX   NCBI_TaxID=70863;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=MR-1;
RX   MEDLINE=22297686; PubMed=12368813; DOI=10.1038/nbt749;
RA   Heidelberg J.F., Paulsen I.T., Nelson K.E., Gaidos E.J., Nelson W.C.,
RA   Read T.D., Eisen J.A., Seshadri R., Ward N.L., Methe B.A.,
RA   Clayton R.A., Meyer T., Tsapin A., Scott J., Beanan M.J.,
RA   Brinkac L.M., Daugherty S.C., DeBoy R.T., Dodson R.J., Durkin A.S.,
RA   Haft D.H., Kolonay J.F., Madupu R., Peterson J.D., Umayam L.A.,
RA   White O., Wolf A.M., Vamathevan J.J., Weidman J.F., Impraim M.,
RA   Lee K., Berry K.J., Lee C., Mueller J., Khouri H.M., Gill J.,
RA   Utterback T.R., McDonald L.A., Feldblyum T.V., Smith H.O.,
RA   Venter J.C., Nealson K.H., Fraser C.M.;
RT   "Genome sequence of the dissimilatory metal ion-reducing bacterium
RT   Shewanella oneidensis.";
RL   Nat. Biotechnol. 20:1118-1123(2002).
DR   EMBL; AE015795; AAN56575.1; -; Genomic_DNA.
DR   TIGR; SO3588; -.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome.
SQ   SEQUENCE   189 AA;  20388 MW;  25E6A0986AF2BFCD CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 189;
  Best Local Similarity   50.0%;  Pred. No. 15;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         14 MGFGMTTILLNIHN 27


RESULT 21
Q87S04_VIBPA
ID   Q87S04_VIBPA PRELIMINARY;      PRT;   196 AA.
AC   Q87S04;
DT   01-JUN-2003 (TrEMBLrel. 24, Created)
DT   01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
DT   01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE   Hypothetical protein VP0620.
GN   OrderedLocusNames=VP0620;
OS   Vibrio parahaemolyticus.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Vibrionales;
OC   Vibrionaceae; Vibrio.
OX   NCBI_TaxID=670;
RN   [1]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC   STRAIN=RIMD 2210633 / Serotype O3:K6;
RX   MEDLINE=22508454; PubMed=12620739; DOI=10.1016/S0140-6736(03)12659-1;
RA   Makino K., Oshima K., Kurokawa K., Yokoyama K., Uda T., Tagomori K.,
RA   Iijima Y., Najima M., Nakano M., Yamashita A., Kubota Y., Kimura S.,
RA   Yasunaga T., Honda T., Shinagawa H., Hattori M., Iida T.;
RT   "Genome sequence of Vibrio parahaemolyticus: a pathogenic mechanism
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RT   distinct from that of V. cholerae.";
RL   Lancet 361:743-749(2003).
DR   EMBL; BA000031; BAC58883.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome; Hypothetical protein.
SQ   SEQUENCE   196 AA;  20930 MW;  8FECFC8E367DDA11 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 196;
  Best Local Similarity   50.0%;  Pred. No. 15;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         14 MGFGMTTILLNIHN 27


RESULT 22
Q6LPH7_PHOPR
ID   Q6LPH7_PHOPR PRELIMINARY;      PRT;   197 AA.
AC   Q6LPH7;
DT   05-JUL-2004 (TrEMBLrel. 27, Created)
DT   05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT   05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE   Putative membrane protein.
GN   Name=STY0009; OrderedLocusNames=PBPRA2415;
OS   Photobacterium profundum (Photobacterium sp. (strain SS9)).
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Vibrionales;
OC   Vibrionaceae; Photobacterium.
OX   NCBI_TaxID=74109;
RN   [1]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RX   PubMed=15746425; DOI=10.1126/science.1103341;
RA   Vezzi A., Campanaro S., D'Angelo M., Simonato F., Vitulo N.,
RA   Lauro F.M., Cestaro A., Malacrida G., Simionati B., Cannata N.,
RA   Romualdi C., Bartlett D.H., Valle G.;
RT   "Life at depth: Photobacterium profundum genome sequence and
RT   expression analysis.";
RL   Science 307:1459-1461(2005).
DR   EMBL; CR378671; CAG20799.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome.
SQ   SEQUENCE   197 AA;  21133 MW;  688D1251A006AFF8 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 197;
  Best Local Similarity   50.0%;  Pred. No. 15;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         14 MGFGMTTILLNIHN 27


RESULT 23
Q8DF09_VIBVU
ID   Q8DF09_VIBVU PRELIMINARY;      PRT;   197 AA.
AC   Q8DF09;
DT   01-MAR-2003 (TrEMBLrel. 23, Created)
DT   01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
DT   01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE   Predicted membrane protein.
GN   OrderedLocusNames=VV10416;
OS   Vibrio vulnificus.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Vibrionales;
OC   Vibrionaceae; Vibrio.
OX   NCBI_TaxID=672;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=CMCP6;
RA   Rhee J.H., Kim S.Y., Chung S.S., Kim J.J., Moon Y.H., Jeong H.,
RA   Choy H.E.;
RT   "Complete genome sequence of Vibrio vulnificus CMCP6.";
RL   Submitted (DEC-2002) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; AE016798; AAO08939.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome.
SQ   SEQUENCE   197 AA;  21066 MW;  E1781C8FF21B08B9 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 197;
  Best Local Similarity   50.0%;  Pred. No. 15;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         14 MGFGMTTILLNIHN 27


RESULT 24
Q9KTW0_VIBCH
ID   Q9KTW0_VIBCH PRELIMINARY;      PRT;   197 AA.
AC   Q9KTW0;
DT   01-OCT-2000 (TrEMBLrel. 15, Created)
DT   01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
DT   01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
DE   Hypothetical protein VC0770.
GN   OrderedLocusNames=VC0770;
OS   Vibrio cholerae.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Vibrionales;
OC   Vibrionaceae; Vibrio.
OX   NCBI_TaxID=666;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=El Tor N16961 / Serotype O1;
RX   MEDLINE=20406833; PubMed=10952301; DOI=10.1038/35020000;
RA   Heidelberg J.F., Eisen J.A., Nelson W.C., Clayton R.A., Gwinn M.L.,
RA   Dodson R.J., Haft D.H., Hickey E.K., Peterson J.D., Umayam L.A.,
RA   Gill S.R., Nelson K.E., Read T.D., Tettelin H., Richardson D.L.,
RA   Ermolaeva M.D., Vamathevan J.J., Bass S., Qin H., Dragoi I.,
RA   Sellers P., McDonald L.A., Utterback T.R., Fleischmann R.D.,
RA   Nierman W.C., White O., Salzberg S.L., Smith H.O., Colwell R.R.,
RA   Mekalanos J.J., Venter J.C., Fraser C.M.;
RT   "DNA sequence of both chromosomes of the cholera pathogen Vibrio
RT   cholerae.";
RL   Nature 406:477-483(2000).
DR   EMBL; AE004162; AAF93935.1; -; Genomic_DNA.
DR   PIR; F82282; F82282.
DR   TIGR; VC0770; -.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome; Hypothetical protein.
SQ   SEQUENCE   197 AA;  21120 MW;  5CD361817EB954E1 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 197;
  Best Local Similarity   50.0%;  Pred. No. 15;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         14 MGFGMTTILLNIHN 27


RESULT 25
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Q7MND9_VIBVY
ID   Q7MND9_VIBVY PRELIMINARY;      PRT;   197 AA.
AC   Q7MND9;
DT   01-MAR-2004 (TrEMBLrel. 26, Created)
DT   01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Predicted membrane protein.
GN   OrderedLocusNames=VV0778;
OS   Vibrio vulnificus (strain YJ016).
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Vibrionales;
OC   Vibrionaceae; Vibrio.
OX   NCBI_TaxID=196600;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RX   PubMed=14656965; DOI=10.1101/gr.1295503;
RA   Chen C.-Y., Wu K.-M., Chang Y.-C., Chang C.-H., Tsai H.-C.,
RA   Liao T.-L., Liu Y.-M., Chen H.-J., Shen A.B.-T., Li J.-C., Su T.-L.,
RA   Shao C.-P., Lee C.-T., Hor L.-I., Tsai S.-F.;
RT   "Comparative genome analysis of Vibrio vulnificus, a marine
RT   pathogen.";
RL   Genome Res. 13:2577-2587(2003).
DR   EMBL; BA000037; BAC93542.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome.
SQ   SEQUENCE   197 AA;  21066 MW;  E1781C8FF21B08B9 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 197;
  Best Local Similarity   50.0%;  Pred. No. 15;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         14 MGFGMTTILLNIHN 27


RESULT 26
Q8PYF9_METMA
ID   Q8PYF9_METMA PRELIMINARY;      PRT;   200 AA.
AC   Q8PYF9;
DT   01-OCT-2002 (TrEMBLrel. 22, Created)
DT   01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
DT   01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
DE   Transcriptional regulator.
GN   OrderedLocusNames=MM0903;
OS   Methanosarcina mazei (Methanosarcina frisia).
OC   Archaea; Euryarchaeota; Methanomicrobia; Methanosarcinales;
OC   Methanosarcinaceae; Methanosarcina.
OX   NCBI_TaxID=2209;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=Goe1 / Go1 / ATCC BAA-199 / DSM 3647 / OCM 88;
RX   MEDLINE=22120827; PubMed=12125824;
RA   Deppenmeier U., Johann A., Hartsch T., Merkl R., Schmitz R.A.,
RA   Martinez-Arias R., Henne A., Wiezer A., Baeumer S., Jacobi C.,
RA   Brueggemann H., Lienard T., Christmann A., Boemecke M., Steckel S.,
RA   Bhattacharyya A., Lykidis A., Overbeek R., Klenk H.-P., Gunsalus R.P.,
RA   Fritz H.-J., Gottschalk G.;
RT   "The genome of Methanosarcina mazei: evidence for lateral gene
RT   transfer between Bacteria and Archaea.";
RL   J. Mol. Microbiol. Biotechnol. 4:453-461(2002).
DR   EMBL; AE013316; AAM30599.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome.
SQ   SEQUENCE   200 AA;  21621 MW;  6F8AB042D9B70FF0 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 200;
  Best Local Similarity   50.0%;  Pred. No. 16;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         29 MGFGMTTVLLNIHN 42


RESULT 27
Q8TIY1_METAC
ID   Q8TIY1_METAC PRELIMINARY;      PRT;   201 AA.
AC   Q8TIY1;
DT   01-JUN-2002 (TrEMBLrel. 21, Created)
DT   01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
DT   01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
DE   Fun34 related protein.
GN   OrderedLocusNames=MA4008;
OS   Methanosarcina acetivorans.
OC   Archaea; Euryarchaeota; Methanomicrobia; Methanosarcinales;
OC   Methanosarcinaceae; Methanosarcina.
OX   NCBI_TaxID=2214;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=C2A / ATCC 35395 / DSM 2834;
RX   MEDLINE=21929760; PubMed=11932238; DOI=10.1101/gr.223902;
RA   Galagan J.E., Nusbaum C., Roy A., Endrizzi M.G., Macdonald P.,
RA   FitzHugh W., Calvo S., Engels R., Smirnov S., Atnoor D., Brown A.,
RA   Allen N., Naylor J., Stange-Thomann N., DeArellano K., Johnson R.,
RA   Linton L., McEwan P., McKernan K., Talamas J., Tirrell A., Ye W.,
RA   Zimmer A., Barber R.D., Cann I., Graham D.E., Grahame D.A., Guss A.M.,
RA   Hedderich R., Ingram-Smith C., Kuettner H.C., Krzycki J.A.,
RA   Leigh J.A., Li W., Liu J., Mukhopadhyay B., Reeve J.N., Smith K.,
RA   Springer T.A., Umayam L.A., White O., White R.H., de Macario E.C.,
RA   Ferry J.G., Jarrell K.F., Jing H., Macario A.J.L., Paulsen I.T.,
RA   Pritchett M., Sowers K.R., Swanson R.V., Zinder S.H., Lander E.,
RA   Metcalf W.W., Birren B.;
RT   "The genome of Methanosarcina acetivorans reveals extensive metabolic
RT   and physiological diversity.";
RL   Genome Res. 12:532-542(2002).
DR   EMBL; AE011113; AAM07358.1; -; Genomic_DNA.
DR   GO; GO:0016020; C:membrane; IEA.
DR   InterPro; IPR000791; Grp1_Fun34_YaaH.
DR   Pfam; PF01184; Grp1_Fun34_YaaH; 1.
DR   ProDom; PD010188; Grp1_Fun34_YaaH; 1.
DR   PROSITE; PS01114; GPR1_FUN34_YAAH; 1.
KW   Complete proteome.
SQ   SEQUENCE   201 AA;  21734 MW;  BF6013C034C0DAFE CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 201;
  Best Local Similarity   50.0%;  Pred. No. 16;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:||    :|:||
Db         29 MGFGMTTVLLNIHN 42


RESULT 28
Q4UYJ0_XANCP
ID   Q4UYJ0_XANCP PRELIMINARY;      PRT;   343 AA.
AC   Q4UYJ0;
DT   13-SEP-2005 (TrEMBLrel. 31, Created)
DT   13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
DT   13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
DE   Transcriptional regulator.
GN   ORFNames=XC_0808;
OS   Xanthomonas campestris pv. campestris str. 8004.
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Xanthomonadales;
OC   Xanthomonadaceae; Xanthomonas.
OX   NCBI_TaxID=314565;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
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RC   STRAIN=8004;
RA   Qian W., Jia Y.-T., Ren S.-X., He Y.-Q., Feng J.-X., Lu L.-F.,
RA   Sun Q.-H., Ying G., Tang D.-J., Wu W., Wang L.-F., Jiang B.-L.,
RA   Zeng S.-Y., Gu W.-Y., Lu G., Rong L., Tian Y.-C., Yao Z.-J., Fu G.,
RA   Chen B.-S., Fang R.-X., Qiang B.-Q., Chen Z., Zhao G.-P., Tang J.-L.,
RA   He C.-Z.;
RL   Submitted (FEB-2005) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; CP000050; AAY47883.1; -; Genomic_DNA.
SQ   SEQUENCE   343 AA;  36846 MW;  A30A81FB7DC8C057 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 343;
  Best Local Similarity   66.7%;  Pred. No. 27;
  Matches    8;  Conservative    1;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLH 13
              || | |||:| |
Db        108 GYSMLLSKLNRH 119


RESULT 29
Q8P5I5_XANCP
ID   Q8P5I5_XANCP PRELIMINARY;      PRT;   343 AA.
AC   Q8P5I5;
DT   01-OCT-2002 (TrEMBLrel. 22, Created)
DT   01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Transcriptional regulator.
GN   Name=cebR;
OS   Xanthomonas campestris (pv. campestris).
OC   Bacteria; Proteobacteria; Gammaproteobacteria; Xanthomonadales;
OC   Xanthomonadaceae; Xanthomonas.
OX   NCBI_TaxID=340;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=ATCC 33913 / NCPPB 528;
RX   MEDLINE=22022145; PubMed=12024217; DOI=10.1038/417459a;
RA   da Silva A.C.R., Ferro J.A., Reinach F.C., Farah C.S., Furlan L.R.,
RA   Quaggio R.B., Monteiro-Vitorello C.B., Van Sluys M.A.,
RA   Almeida N.F. Jr., Alves L.M.C., do Amaral A.M., Bertolini M.C.,
RA   Camargo L.E.A., Camarotte G., Cannavan F., Cardozo J., Chambergo F.,
RA   Ciapina L.P., Cicarelli R.M.B., Coutinho L.L., Cursino-Santos J.R.,
RA   El-Dorry H., Faria J.B., Ferreira A.J.S., Ferreira R.C.C.,
RA   Ferro M.I.T., Formighieri E.F., Franco M.C., Greggio C.C., Gruber A.,
RA   Katsuyama A.M., Kishi L.T., Leite R.P., Lemos E.G.M., Lemos M.V.F.,
RA   Locali E.C., Machado M.A., Madeira A.M.B.N., Martinez-Rossi N.M.,
RA   Martins E.C., Meidanis J., Menck C.F.M., Miyaki C.Y., Moon D.H.,
RA   Moreira L.M., Novo M.T.M., Okura V.K., Oliveira M.C., Oliveira V.R.,
RA   Pereira H.A., Rossi A., Sena J.A.D., Silva C., de Souza R.F.,
RA   Spinola L.A.F., Takita M.A., Tamura R.E., Teixeira E.C., Tezza R.I.D.,
RA   Trindade dos Santos M., Truffi D., Tsai S.M., White F.F.,
RA   Setubal J.C., Kitajima J.P.;
RT   "Comparison of the genomes of two Xanthomonas pathogens with differing
RT   host specificities.";
RL   Nature 417:459-463(2002).
DR   EMBL; AE012454; AAM42626.1; -; Genomic_DNA.
DR   HSSP; P03023; 1CJG.
DR   GO; GO:0005622; C:intracellular; IEA.
DR   GO; GO:0003700; F:transcription factor activity; IEA.
DR   GO; GO:0006355; P:regulation of transcription, DNA-dependent; IEA.
DR   InterPro; IPR000843; HTH_LacI.
DR   InterPro; IPR001761; PeriplaBP/LacI.
DR   Pfam; PF00356; LacI; 1.
DR   Pfam; PF00532; Peripla_BP_1; 1.
DR   SMART; SM00354; HTH_LACI; 1.
DR   PROSITE; PS00356; HTH_LACI_1; 1.
DR   PROSITE; PS50932; HTH_LACI_2; 1.
KW   Complete proteome.
SQ   SEQUENCE   343 AA;  36846 MW;  A30A81FB7DC8C057 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 343;
  Best Local Similarity   66.7%;  Pred. No. 27;
  Matches    8;  Conservative    1;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLH 13
              || | |||:| |
Db        108 GYSMLLSKLNRH 119


RESULT 30
Q6MST2_MYCMS
ID   Q6MST2_MYCMS PRELIMINARY;      PRT;   362 AA.
AC   Q6MST2;
DT   05-JUL-2004 (TrEMBLrel. 27, Created)
DT   05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT   05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE   Probable nicotinic phosphoribosyltransferase (EC 2.4.2.11).
GN   Name=pncB; OrderedLocusNames=MSC_0687;
OS   Mycoplasma mycoides (subsp. mycoides SC).
OC   Bacteria; Firmicutes; Mollicutes; Mycoplasmataceae; Mycoplasma.
OX   NCBI_TaxID=44101;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=PG1;
RX   PubMed=14762060; DOI=10.1101/gr.1673304;
RA   Westberg J., Persson A., Holmberg A., Goesmann A., Lundeberg J.,
RA   Johansson K.-E., Pettersson B., Uhlen M.;
RT   "The genome sequence of Mycoplasma mycoides subsp. mycoides SC type
RT   strain PG1T, the causative agent of contagious bovine pleuropneumonia
RT   (CBPP).";
RL   Genome Res. 14:221-227(2004).
DR   EMBL; BX842644; CAE77306.1; -; Genomic_DNA.
DR   GO; GO:0004516; F:nicotinate phosphoribosyltransferase activity; IEA.
KW   Complete proteome.
SQ   SEQUENCE   362 AA;  41160 MW;  8121BF091281DFB9 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 362;
  Best Local Similarity   66.7%;  Pred. No. 28;
  Matches    8;  Conservative    2;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLH 13
              | | ||:|||:|
Db        321 GVGSALAKINIH 332


RESULT 31
Q4H6L8_9DEIO
ID   Q4H6L8_9DEIO PRELIMINARY;      PRT;   501 AA.
AC   Q4H6L8;
DT   13-SEP-2005 (TrEMBLrel. 31, Created)
DT   13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
DT   13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
DE   Similar to metal-dependent hydrolase with the TIM-barrel fold.
GN   ORFNames=DgeoDRAFT_0101;
OS   Deinococcus geothermalis DSM 11300.
OC   Bacteria; Deinococcus-Thermus; Deinococci; Deinococcales;
OC   Deinococcaceae; Deinococcus.
OX   NCBI_TaxID=319795;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=DSM 11300;
RG   US DOE Joint Genome Institute (JGI-PGF);
RA   Copeland A., Lucas S., Lapidus A., Barry K., Detter C., Glavina T.,
RA   Hammon N., Israni S., Pitluck S., Richardson P.;
RT   "Sequencing of the draft genome assembly of Deinococcus geothermalis
RT   DSM 11300.";
RL   Submitted (MAY-2005) to the EMBL/GenBank/DDBJ databases.
RN   [2]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=DSM 11300;
RG   US DOE Joint Genome Institute (JGI-ORNL);
RA   Larimer F., Land M.;
RT   "Annotation of the draft genome assembly of Deinococcus geothermalis
RT   DSM 11300.";
RL   Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases.
CC   -!- CAUTION: The sequence shown here is derived from an
CC       EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
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CC       preliminary data.
DR   EMBL; AAHE01000013; EAL81904.1; -; Genomic_DNA.
KW   Hydrolase.
SQ   SEQUENCE   501 AA;  53099 MW;  13CFBB50B9CC64AB CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 501;
  Best Local Similarity   53.8%;  Pred. No. 39;
  Matches    7;  Conservative    4;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLH 13
              : || :||::|||
Db         72 VAYGFSLSQLNLH 84


RESULT 32
Q4RYL2_TETNG
ID   Q4RYL2_TETNG PRELIMINARY;      PRT;   694 AA.
AC   Q4RYL2;
DT   13-SEP-2005 (TrEMBLrel. 31, Created)
DT   13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
DT   13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
DE   Chromosome 3 SCAF14975, whole genome shotgun sequence.
DE   (Fragment).
GN   ORFNames=GSTENG00026906001;
OS   Tetraodon nigroviridis (Green puffer).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
OC   Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
OC   Tetradontoidea; Tetraodontidae; Tetraodon.
OX   NCBI_TaxID=99883;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RA   Jaillon O., Aury J.M., Brunet F., Petit J.L., Stange-Thomann N.,
RA   Mauceli E., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,
RA   Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,
RA   Dasilva C., Salanoubat M., Levy M., Boudet N., Castellano S.,
RA   Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
RA   Biemont C., Skalli Z., Cattolico L., Poulain J., De Berardinis V.,
RA   Cruaud C., Duprat S., Brottier P., Coutanceau J.P., Gouzy J.,
RA   Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,
RA   Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,
RA   Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
RA   Laudet V., Schachter V., Quetier F., Saurin W., Scarpelli C.,
RA   Wincker P., Lander E.S., Weissenbach J., Roest Crollius H.;
RT   "Genome duplication in the teleost fish Tetraodon nigroviridis reveals
RT   the early vertebrate proto-karyotype.";
RL   Nature 431:946-957(2004).
RN   [2]
RP   NUCLEOTIDE SEQUENCE.
RG   Genoscope; Whitehead Institute Centre for Genome Research;
RL   Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.
CC   -!- CAUTION: The sequence shown here is derived from an
CC       EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
CC       preliminary data.
DR   EMBL; CAAE01014975; CAG06520.1; -; Genomic_DNA.
FT   NON_TER       1      1
FT   NON_TER     694    694
SQ   SEQUENCE   694 AA;  77492 MW;  FDBB74F7D7F17FE7 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 694;
  Best Local Similarity   72.7%;  Pred. No. 54;
  Matches    8;  Conservative    3;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::||||:|||
Db        496 GISLSKISLHN 506


RESULT 33
Q899T2_CLOTE
ID   Q899T2_CLOTE PRELIMINARY;      PRT;   887 AA.
AC   Q899T2;
DT   01-JUN-2003 (TrEMBLrel. 24, Created)
DT   01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
DT   01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE   Conserved membrane protein.
GN   OrderedLocusNames=CTC00086;
OS   Clostridium tetani.
OC   Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;
OC   Clostridium.
OX   NCBI_TaxID=1513;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=Massachusetts / E88;
RX   MEDLINE=22457253; PubMed=12552129; DOI=10.1073/pnas.0335853100;
RA   Brueggemann H., Baeumer S., Fricke W.F., Wiezer A., Liesegang H.,
RA   Decker I., Herzberg C., Martinez-Arias R., Merkl R., Henne A.,
RA   Gottschalk G.;
RT   "The genome sequence of Clostridium tetani, the causative agent of
RT   tetanus disease.";
RL   Proc. Natl. Acad. Sci. U.S.A. 100:1316-1321(2003).
DR   EMBL; AE015936; AAO34739.1; -; Genomic_DNA.
DR   GO; GO:0016021; C:integral to membrane; IEA.
DR   InterPro; IPR005372; UPF0182.
DR   Pfam; PF03699; UPF0182; 1.
KW   Complete proteome.
SQ   SEQUENCE   887 AA;  103741 MW;  A92998D2EA3CB7D7 CRC64;

  Query Match             59.5%;  Score 44;  DB 2;  Length 887;
  Best Local Similarity   70.0%;  Pred. No. 68;
  Matches    7;  Conservative    3;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKIN 11
              |||:|:||:|
Db        417 GYGVAMSKVN 426


RESULT 34
RS24_AERPE
ID   RS24_AERPE     STANDARD;      PRT;   102 AA.
AC   Q9YCY0;
DT   30-MAY-2000 (Rel. 39, Created)
DT   30-MAY-2000 (Rel. 39, Last sequence update)
DT   10-MAY-2005 (Rel. 47, Last annotation update)
DE   30S ribosomal protein S24e.
GN   Name=rps24e; OrderedLocusNames=APE1132;
OS   Aeropyrum pernix.
OC   Archaea; Crenarchaeota; Thermoprotei; Desulfurococcales;
OC   Desulfurococcaceae; Aeropyrum.
OX   NCBI_TaxID=56636;
RN   [1]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC   STRAIN=K1;
RX   MEDLINE=99310339; PubMed=10382966;
RA   Kawarabayasi Y., Hino Y., Horikawa H., Yamazaki S., Haikawa Y.,
RA   Jin-no K., Takahashi M., Sekine M., Baba S.-I., Ankai A., Kosugi H.,
RA   Hosoyama A., Fukui S., Nagai Y., Nishijima K., Nakazawa H.,
RA   Takamiya M., Masuda S., Funahashi T., Tanaka T., Kudoh Y.,
RA   Yamazaki J., Kushida N., Oguchi A., Aoki K.-I., Kubota K.,
RA   Nakamura Y., Nomura N., Sako Y., Kikuchi H.;
RT   "Complete genome sequence of an aerobic hyper-thermophilic
RT   crenarchaeon, Aeropyrum pernix K1.";
RL   DNA Res. 6:83-101(1999).
CC   -!- SIMILARITY: Belongs to the ribosomal protein S24e family.
CC   ---------------------------------------------------------------------
CC   This Swiss-Prot entry is copyright. It is produced through a collaboration
CC   between the Swiss Institute of Bioinformatics and the EMBL outstation -
CC   the European Bioinformatics Institute. There are no restrictions on its
CC   use as long as its content is in no way modified and this statement is not
CC   removed.
CC   ---------------------------------------------------------------------
DR   EMBL; BA000002; BAA80117.1; ALT_INIT; Genomic_DNA.
DR   HAMAP; MF_00545; -; 1.
DR   InterPro; IPR001976; Ribosomal_S24E.
DR   PANTHER; PTHR10496; Ribosomal_S24E; 1.
DR   Pfam; PF01282; Ribosomal_S24e; 1.
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DR   ProDom; PD006052; Ribosomal_S24E; 1.
DR   PROSITE; PS00529; RIBOSOMAL_S24E; 1.
KW   Complete proteome; Ribonucleoprotein; Ribosomal protein.
SQ   SEQUENCE   102 AA;  11858 MW;  DEAA205AAFBD8066 CRC64;

  Query Match             58.1%;  Score 43;  DB 1;  Length 102;
  Best Local Similarity   58.3%;  Pred. No. 12;
  Matches    7;  Conservative    2;  Mismatches    3;  Indels    0;  Gaps    0;

Qy          2 GYGMALSKINLH 13
              |||  |||: :|
Db         60 GYGAGLSKVRVH 71


RESULT 35
Q4NQB2_9DELT
ID   Q4NQB2_9DELT PRELIMINARY;      PRT;   214 AA.
AC   Q4NQB2;
DT   13-SEP-2005 (TrEMBLrel. 31, Created)
DT   13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
DT   13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
DE   GPR1/FUN34/yaaH.
GN   ORFNames=AdehDRAFT_0850;
OS   Anaeromyxobacter dehalogenans 2CP-C.
OC   Bacteria; Proteobacteria; Deltaproteobacteria; Myxococcales;
OC   Cystobacterineae; Myxococcaceae; Anaeromyxobacter.
OX   NCBI_TaxID=290397;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=2CP-C;
RG   US DOE Joint Genome Institute (JGI-PGF);
RA   Copeland A., Lucas S., Lapidus A., Barry K., Detter C., Glavina T.,
RA   Hammon N., Israni S., Pitluck S., Richardson P.;
RT   "Sequencing of the draft genome assembly of Anaeromyxobacter
RT   dehalogenans 2CP-C.";
RL   Submitted (MAY-2005) to the EMBL/GenBank/DDBJ databases.
RN   [2]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=2CP-C;
RG   US DOE Joint Genome Institute (JGI-ORNL);
RA   Larimer F., Land M.;
RT   "Annotation of the draft genome assembly of Anaeromyxobacter
RT   dehalogenans 2CP-C.";
RL   Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases.
CC   -!- CAUTION: The sequence shown here is derived from an
CC       EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
CC       preliminary data.
DR   EMBL; AAHD01000036; EAL77772.1; -; Genomic_DNA.
SQ   SEQUENCE   214 AA;  22798 MW;  9C3059FF2E547170 CRC64;

  Query Match             58.1%;  Score 43;  DB 2;  Length 214;
  Best Local Similarity   50.0%;  Pred. No. 25;
  Matches    7;  Conservative    3;  Mismatches    4;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLHN 14
              ||:|:    :||||
Db         42 MGFGLTTVLLNLHN 55


RESULT 36
Q4IKZ0_GIBZE
ID   Q4IKZ0_GIBZE PRELIMINARY;      PRT;   541 AA.
AC   Q4IKZ0;
DT   13-SEP-2005 (TrEMBLrel. 31, Created)
DT   13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
DT   13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
DE   Hypothetical protein.
GN   ORFNames=FG02118.1;
OS   Gibberella zeae PH-1.
OC   Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes;
OC   Hypocreomycetidae; Hypocreales; Nectriaceae; Gibberella.
OX   NCBI_TaxID=229533;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   STRAIN=PH-1;
RA   Birren B., Nusbaum C., Abouelleil A., Allen N., Anderson S.,
RA   Arachchi H.M., Barna N., Bastien V., Bloom T., Boguslavkiy L.,
RA   Boukhgalter B., Butler J., Calvo S.E., Camarata J., Chang J.,
RA   Choepel Y., Collymore A., Cook A., Cooke P., Corum B., DeArellano K.,
RA   Diaz J.S., Dodge S., Dooley K., Dorris L., Elkins T., Engels R.,
RA   Erickson J., Faro S., Ferreira P., FitzGerald M., Gage D., Galagan J.,
RA   Gardyna S., Gnerre S., Graham L., Grand-Pierre N., Hafez N.,
RA   Hagopian D., Hagos B., Hall J., Horton L., Hulme W., Iliev I.,
RA   Jaffe D., Johnson R., Jones C., Kamal M., Kamat A., Karatas A.,
RA   Kells C., Landers T., Levine R., Lindblad-Toh K., Liu G., Lui A.,
RA   Ma L.-J., Mabbitt R., MacLean C., Macdonald P., Major J., Manning J.,
RA   Matthews C., Mauceli E., McCarthy M., Meldrim J., Meneus L.,
RA   Mihova T., Mlenga V., Murphy T., Naylor J., Nguyen C., Nicol R.,
RA   Nielsen C.B., Norbu C., O'Connor T., O'Donnell P., O'Neil D.,
RA   Oliver J., Peterson K., Phunkhang P., Pierre N., Purcell S.,
RA   Rachupka A., Ramasamy U., Raymond C., Retta R., Rise C., Rogov P.,
RA   Roman J., Schauer S., Schupback R., Seaman S., Severy P., Smirnov S.,
RA   Smith C., Spencer B., Stange-Thomann N., Stojanovic N., Stubbs M.,
RA   Talamas J., Tesfaye S., Theodore J., Topham K., Travers M.,
RA   Vassiliev H., Venkataraman V.S., Viel R., Vo A., Wang S., Wilson B.,
RA   Wu X., Wyman D., Young G., Zainoun J., Zembek L., Zimmer A., Zody M.,
RA   Lander E.;
RT   "Fusarium graminearum genome sequence.";
RL   Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.
CC   -!- CAUTION: The sequence shown here is derived from an
CC       EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
CC       preliminary data.
DR   EMBL; AACM01000111; EAA69749.1; -; Genomic_DNA.
KW   Hypothetical protein.
SQ   SEQUENCE   541 AA;  60552 MW;  D9EEFE69CD5AFDBB CRC64;

  Query Match             56.8%;  Score 42;  DB 2;  Length 541;
  Best Local Similarity   53.8%;  Pred. No. 97;
  Matches    7;  Conservative    4;  Mismatches    2;  Indels    0;  Gaps    0;

Qy          1 MGYGMALSKINLH 13
              :|||: ||: |:|
Db        153 LGYGLLLSEGNVH 165


RESULT 37
SLIK1_HUMAN
ID   SLIK1_HUMAN     STANDARD;      PRT;   696 AA.
AC   Q96PX8; Q96SF9;
DT   05-JUL-2004 (Rel. 44, Created)
DT   05-JUL-2004 (Rel. 44, Last sequence update)
DT   13-SEP-2005 (Rel. 48, Last annotation update)
DE   SLIT and NTRK-like protein 1 precursor.
GN   Name=SLITRK1; Synonyms=KIAA1910, LRCC12; ORFNames=UNQ233/PRO266;
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
OC   Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
RC   TISSUE=Brain;
RX   MEDLINE=21456161; PubMed=11572484;
RA   Nagase T., Kikuno R., Ohara O.;
RT   "Prediction of the coding sequences of unidentified human genes. XXI.
RT   The complete sequences of 60 new cDNA clones from brain which code for
RT   large proteins.";
RL   DNA Res. 8:179-187(2001).
RN   [2]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
RX   MEDLINE=22887296; PubMed=12975309; DOI=10.1101/gr.1293003;
RA   Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D.T., Brush J.,
RA   Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,
RA   Eaton D., Foster J.S., Grimaldi C., Gu Q., Hass P.E., Heldens S.,
RA   Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,
RA   Lewis L., Liao D., Mark M.R., Robbie E., Sanchez C., Schoenfeld J.,
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RA   Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,
RA   Vandlen R.L., Watanabe C., Wieand D., Woods K., Xie M.-H.,
RA   Yansura D.G., Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A.D.,
RA   Wood W.I., Godowski P.J., Gray A.M.;
RT   "The secreted protein discovery initiative (SPDI), a large-scale
RT   effort to identify novel human secreted and transmembrane proteins: a
RT   bioinformatics assessment.";
RL   Genome Res. 13:2265-2270(2003).
RN   [3]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RX   PubMed=15057823; DOI=10.1038/nature02379;
RA   Dunham A., Matthews L.H., Burton J., Ashurst J.L., Howe K.L.,
RA   Ashcroft K.J., Beare D.M., Burford D.C., Hunt S.E.,
RA   Griffiths-Jones S., Jones M.C., Keenan S.J., Oliver K., Scott C.E.,
RA   Ainscough R., Almeida J.P., Ambrose K.D., Andrews D.T.,
RA   Ashwell R.I.S., Babbage A.K., Bagguley C.L., Bailey J., Bannerjee R.,
RA   Barlow K.F., Bates K., Beasley H., Bird C.P., Bray-Allen S.,
RA   Brown A.J., Brown J.Y., Burrill W., Carder C., Carter N.P.,
RA   Chapman J.C., Clamp M.E., Clark S.Y., Clarke G., Clee C.M.,
RA   Clegg S.C., Cobley V., Collins J.E., Corby N., Coville G.J.,
RA   Deloukas P., Dhami P., Dunham I., Dunn M., Earthrowl M.E.,
RA   Ellington A.G., Faulkner L., Frankish A.G., Frankland J., French L.,
RA   Garner P., Garnett J., Gilbert J.G.R., Gilson C.J., Ghori J.,
RA   Grafham D.V., Gribble S.M., Griffiths C., Hall R.E., Hammond S.,
RA   Harley J.L., Hart E.A., Heath P.D., Howden P.J., Huckle E.J.,
RA   Hunt P.J., Hunt A.R., Johnson C., Johnson D., Kay M., Kimberley A.M.,
RA   King A., Laird G.K., Langford C.J., Lawlor S., Leongamornlert D.A.,
RA   Lloyd D.M., Lloyd C., Loveland J.E., Lovell J., Martin S.,
RA   Mashreghi-Mohammadi M., McLaren S.J., McMurray A., Milne S.,
RA   Moore M.J.F., Nickerson T., Palmer S.A., Pearce A.V., Peck A.I.,
RA   Pelan S., Phillimore B., Porter K.M., Rice C.M., Searle S.,
RA   Sehra H.K., Shownkeen R., Skuce C.D., Smith M., Steward C.A.,
RA   Sycamore N., Tester J., Thomas D.W., Tracey A., Tromans A., Tubby B.,
RA   Wall M., Wallis J.M., West A.P., Whitehead S.L., Willey D.L.,
RA   Wilming L., Wray P.W., Wright M.W., Young L., Coulson A., Durbin R.,
RA   Hubbard T., Sulston J.E., Beck S., Bentley D.R., Rogers J., Ross M.T.;
RT   "The DNA sequence and analysis of human chromosome 13.";
RL   Nature 428:522-528(2004).
RN   [4]
RP   IDENTIFICATION, AND TISSUE SPECIFICITY.
RC   TISSUE=Brain, and Brain tumor;
RX   PubMed=14557068; DOI=10.1016/S0378-1119(03)00715-7;
RA   Aruga J., Yokota N., Mikoshiba K.;
RT   "Human SLITRK family genes: genomic organization and expression
RT   profiling in normal brain and brain tumor tissue.";
RL   Gene 315:87-94(2003).
CC   -!- FUNCTION: Enhances neurite outgrowth (By similarity).
CC   -!- SUBCELLULAR LOCATION: Membrane-bound (Potential).
CC   -!- TISSUE SPECIFICITY: Expressed predominantly in the frontal lobe of
CC       the cerebral cortex of the brain. Also expressed in some
CC       astrocytic brain tumors such as astrocytomas, oligodendrogliomas,
CC       gliobastomas, gangliogliomas and primitive neuroectodermal tumors.
CC   -!- SIMILARITY: Belongs to the SLITRK family.
CC   -!- SIMILARITY: Contains 13 LRR (leucine-rich) repeats.
CC   ----------------------------------------------------------------------
CC   This Swiss-Prot entry is copyright. It is produced through a collaboration
CC   between the Swiss Institute of Bioinformatics and the EMBL outstation -
CC   the European Bioinformatics Institute. There are no restrictions on its
CC   use as long as its content is in no way modified and this statement is not
CC   removed.
CC   ----------------------------------------------------------------------
DR   EMBL; AB067497; BAB67803.1; ALT_INIT; mRNA.
DR   EMBL; AY358289; AAQ88656.1; -; mRNA.
DR   EMBL; AL355481; CAC37488.1; -; Genomic_DNA.
DR   Ensembl; ENSG00000178235; Homo sapiens.
DR   HGNC; HGNC:20297; SLITRK1.
DR   InterPro; IPR001611; LRR.
DR   InterPro; IPR000483; LRR_Cterm.
DR   InterPro; IPR003591; LRR_typ.
DR   Pfam; PF00560; LRR_1; 8.
DR   PRINTS; PR00019; LEURICHRPT.
DR   SMART; SM00369; LRR_TYP; 11.
DR   SMART; SM00082; LRRCT; 2.
KW   Leucine-rich repeat; Repeat; Signal; Transmembrane.
FT   SIGNAL        1     17       Potential.
FT   CHAIN        18    696       SLIT and NTRK-like protein 1.
FT   TRANSMEM    623    643       Potential.
FT   REPEAT       57     80       LRR 1.
FT   REPEAT       81    104       LRR 2.
FT   REPEAT      105    128       LRR 3.
FT   REPEAT      130    152       LRR 4.
FT   REPEAT      153    176       LRR 5.
FT   REPEAT      178    200       LRR 6.
FT   REPEAT      214    237       LRR 7.
FT   REPEAT      374    397       LRR 8.
FT   REPEAT      399    421       LRR 9.
FT   REPEAT      423    445       LRR 10.
FT   REPEAT      446    469       LRR 11.
FT   REPEAT      470    493       LRR 12.
FT   REPEAT      495    517       LRR 13.
SQ   SEQUENCE   696 AA;  77735 MW;  E0E9ACEDE0F0ACEC CRC64;

  Query Match             56.8%;  Score 42;  DB 1;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.2e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 38
SLIK1_MOUSE
ID   SLIK1_MOUSE    STANDARD;      PRT;   696 AA.
AC   Q810C1; Q9CXL0;
DT   05-JUL-2004 (Rel. 44, Created)
DT   05-JUL-2004 (Rel. 44, Last sequence update)
DT   10-MAY-2005 (Rel. 47, Last annotation update)
DE   SLIT and NTRK-like protein 1 precursor.
GN   Name=Slitrk1; Synonyms=Sltk1;
OS   Mus musculus (Mouse).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
OC   Muroidea; Muridae; Murinae; Mus.
OX   NCBI_TaxID=10090;
RN   [1]
RP   NUCLEOTIDE SEQUENCE, FUNCTION, TISSUE SPECIFICITY, AND DEVELOPMENTAL
RP   STAGE.
RX   PubMed=14550773; DOI=10.1016/S1044-7431(03)00129-5;
RA   Aruga J., Mikoshiba K.;
RT   "Identification and characterization of Slitrk, a novel neuronal
RT   transmembrane protein family controlling neurite outgrowth.";
RL   Mol. Cell. Neurosci. 24:117-129(2003).
RN   [2]
RP   NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA] OF 369-696.
RC   STRAIN=C57BL/6J; TISSUE=Head;
RX   MEDLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;
RA   Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,
RA   Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,
RA   Yagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori T.,
RA   Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
RA   Schriml L.M., Kanapin A., Matsuda H., Batalov S., Beisel K.W.,
RA   Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA   Dalla E., Dragani T.A., Fletcher C.F., Forrest A., Frazer K.S.,
RA   Gaasterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,
RA   Grimmond S., Gustincich S., Hirokawa N., Jackson I.J., Jarvis E.D.,
RA   Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
RA   Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
RA   Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,
RA   Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,
RA   Petrovsky N., Pillai R., Pontius J.U., Qi D., Ramachandran S.,
RA   Ravasi T., Reed J.C., Reed D.J., Reid J., Ring B.Z., Ringwald M.,
RA   Sandelin A., Schneider C., Semple C.A., Setou M., Shimada K.,
RA   Sultana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M.,
RA   Verardo R., Wagner L., Wahlestedt C., Wang Y., Watanabe Y., Wells C.,
RA   Wilming L.G., Wynshaw-Boris A., Yanagisawa M., Yang I., Yang L.,
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RA   Yuan Z., Zavolan M., Zhu Y., Zimmer A., Carninci P., Hayatsu N.,
RA   Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato K.,
RA   Shiraki T., Waki K., Kawai J., Aizawa K., Arakawa T., Fukuda S.,
RA   Hara A., Hashizume W., Imotani K., Ishii Y., Itoh M., Kagawa I.,
RA   Miyazaki A., Sakai K., Sasaki D., Shibata K., Shinagawa A.,
RA   Yasunishi A., Yoshino M., Waterston R., Lander E.S., Rogers J.,
RA   Birney E., Hayashizaki Y.;
RT   "Analysis of the mouse transcriptome based on functional annotation of
RT   60,770 full-length cDNAs.";
RL   Nature 420:563-573(2002).
CC   -!- FUNCTION: Enhances neurite outgrowth.
CC   -!- SUBCELLULAR LOCATION: Membrane-bound (Potential).
CC   -!- TISSUE SPECIFICITY: In the adult, significant expression is
CC       detected only in the brain. Broadly expressed in embryonic brain
CC       with highest expression in subventricular zone, subplate, cortical
CC       plate, pyramidal cell layer of hippocampus, thalamus and
CC       hypothalamus where levels are highest in ventromedial hypothalamus
CC       and medial part of periacqueductal gray matter. Also expressed in
CC       mantle layer of spinal cord and in lateral and medial motor
CC       columns.
CC   -!- DEVELOPMENTAL STAGE: In the embryo, expressed from day 10-12 and
CC       continues through later gestational development and into
CC       adulthood.
CC   -!- SIMILARITY: Belongs to the SLITRK family.
CC   -!- SIMILARITY: Contains 13 LRR (leucine-rich) repeats.
CC   ----------------------------------------------------------------------
CC   This Swiss-Prot entry is copyright. It is produced through a collaboration
CC   between the Swiss Institute of Bioinformatics and the EMBL outstation -
CC   the European Bioinformatics Institute. There are no restrictions on its
CC   use as long as its content is in no way modified and this statement is not
CC   removed.
CC   ----------------------------------------------------------------------
DR   EMBL; AB097570; BAC67204.1; -; Genomic_DNA.
DR   EMBL; AK014285; BAB29244.2; -; mRNA.
DR   HSSP; Q9BZR6; 1OZN.
DR   Ensembl; ENSMUSG00000052049; Mus musculus.
DR   MGI; MGI:2679446; Slitrk1.
DR   GO; GO:0007409; P:axonogenesis; IDA.
DR   InterPro; IPR001611; LRR.
DR   InterPro; IPR000483; LRR_Cterm.
DR   InterPro; IPR000372; LRR_Nterm.
DR   InterPro; IPR003591; LRR_typ.
DR   Pfam; PF00560; LRR_1; 8.
DR   PRINTS; PR00019; LEURICHRPT.
DR   SMART; SM00369; LRR_TYP; 2.
DR   SMART; SM00082; LRRCT; 2.
DR   SMART; SM00013; LRRNT; 1.
KW   Leucine-rich repeat; Repeat; Signal; Transmembrane.
FT   SIGNAL        1     17       Potential.
FT   CHAIN        18    696       SLIT and NTRK-like protein 1.
FT   TRANSMEM    623    643       Potential.
FT   REPEAT       57     80       LRR 1.
FT   REPEAT       81    104       LRR 2.
FT   REPEAT      105    128       LRR 3.
FT   REPEAT      130    152       LRR 4.
FT   REPEAT      153    176       LRR 5.
FT   REPEAT      178    200       LRR 6.
FT   REPEAT      214    237       LRR 7.
FT   REPEAT      374    397       LRR 8.
FT   REPEAT      399    421       LRR 9.
FT   REPEAT      423    445       LRR 10.
FT   REPEAT      446    469       LRR 11.
FT   REPEAT      470    493       LRR 12.
FT   REPEAT      495    517       LRR 13.
SQ   SEQUENCE   696 AA;  77817 MW;  B3F23AF1A28B9E1B CRC64;

  Query Match             56.8%;  Score 42;  DB 1;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.2e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503
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RESULT 39
Q5U5I6_HUMAN
ID   Q5U5I6_HUMAN PRELIMINARY;      PRT;   696 AA.
AC   Q5U5I6;
DT   01-FEB-2005 (TrEMBLrel. 29, Created)
DT   01-FEB-2005 (TrEMBLrel. 29, Last sequence update)
DT   01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
DE   Slit and trk like 1 protein.
GN   Name=SLITRK1;
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
OC   Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   TISSUE=Brain;
RX   MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA   Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA   Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA   Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA   Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA   Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA   Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA   Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA   Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA   Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA   Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA   Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA   Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA   Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA   Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA   Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA   Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
RA   Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
RT   "Generation and initial analysis of more than 15,000 full-length human
RT   and mouse cDNA sequences.";
RL   Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN   [2]
RP   NUCLEOTIDE SEQUENCE.
RC   TISSUE=Brain;
RA   Director MGC Project;
RL   Submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; BC051738; AAH51738.1; -; mRNA.
DR   InterPro; IPR001611; LRR.
DR   InterPro; IPR000483; LRR_Cterm.
DR   InterPro; IPR003591; LRR_typ.
DR   Pfam; PF00560; LRR_1; 8.
DR   PRINTS; PR00019; LEURICHRPT.
DR   SMART; SM00369; LRR_TYP; 11.
DR   SMART; SM00082; LRRCT; 2.
KW   Leucine-rich repeat; Repeat.
SQ   SEQUENCE   696 AA;  77721 MW;  FC82AFCCD2E0ADBD CRC64;

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.2e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


RESULT 40
Q5RAC4_PONPY
ID   Q5RAC4_PONPY PRELIMINARY;      PRT;   696 AA.
AC   Q5RAC4;
DT   01-FEB-2005 (TrEMBLrel. 29, Created)
DT   01-FEB-2005 (TrEMBLrel. 29, Last sequence update)
DT   01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
DE   Hypothetical protein DKFZp459G0529.
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GN   Name=DKFZp459G0529;
OS   Pongo pygmaeus (Orangutan).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
OC   Pongo.
OX   NCBI_TaxID=9600;
RN   [1]
RP   NUCLEOTIDE SEQUENCE.
RC   TISSUE=Cortex;
RG   The German cDNA Consortium;
RA   Bloecker H., Boecher M., Brandt P., Mewes H.W., Weil B., Amid C.,
RA   Osanger A., Fobo G., Han M., Wiemann S.;
RL   Submitted (NOV-2004) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; CR859094; CAH91286.1; -; mRNA.
DR   InterPro; IPR001611; LRR.
DR   InterPro; IPR000483; LRR_Cterm.
DR   InterPro; IPR003591; LRR_typ.
DR   Pfam; PF00560; LRR_1; 8.
DR   PRINTS; PR00019; LEURICHRPT.
DR   SMART; SM00369; LRR_TYP; 11.
DR   SMART; SM00082; LRRCT; 2.
KW   Hypothetical protein; Leucine-rich repeat; Repeat.
SQ   SEQUENCE   696 AA;  77699 MW;  7AF48D3844D97CBC CRC64;

  Query Match             56.8%;  Score 42;  DB 2;  Length 696;
  Best Local Similarity   63.6%;  Pred. No. 1.2e+02;
  Matches    7;  Conservative    4;  Mismatches    0;  Indels    0;  Gaps    0;

Qy          4 GMALSKINLHN 14
              |::|||::|||
Db        493 GVSLSKLSLHN 503


Search completed: May 13, 2006, 08:13:59
Job time : 232 secs
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